

# ToxCast / Tox21 Data Overview

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# A Problem Driving CompTox Development

- ~1,000 chemicals – most people are often exposed to
- ~10,000 chemicals – many people are exposed to at least occasionally
- ~100,000 chemicals – some people are exposed to at least occasionally
  
- Some of these chemicals / exposures are responsible for
  - Cancers
  - Birth defects
  - Reproductive disorders
  - Both human and ecological species
  
- Can we prioritize chemicals for further testing without knowing everything?

## ToxCast Projects

- ToxCast / Tox21 is a large-scale *in vitro* screening program
  - ToxCast: 1,000 chemicals in ~500 assays
  - Tox21: 10,000 chemicals in ~50 assays (drugs+EPA+NTP chemicals)
- Reverse toxicokinetics (RTK) approach for estimating maximum “no bioactivity” dose estimate
- Input for Endocrine Screening Prioritization and other programs
- Models for Prioritization of Targeted Testing
  - Test examples have known MOA, activity
  - Goal is to qualify *in vitro* assays for use in prioritizing testing of untested compounds

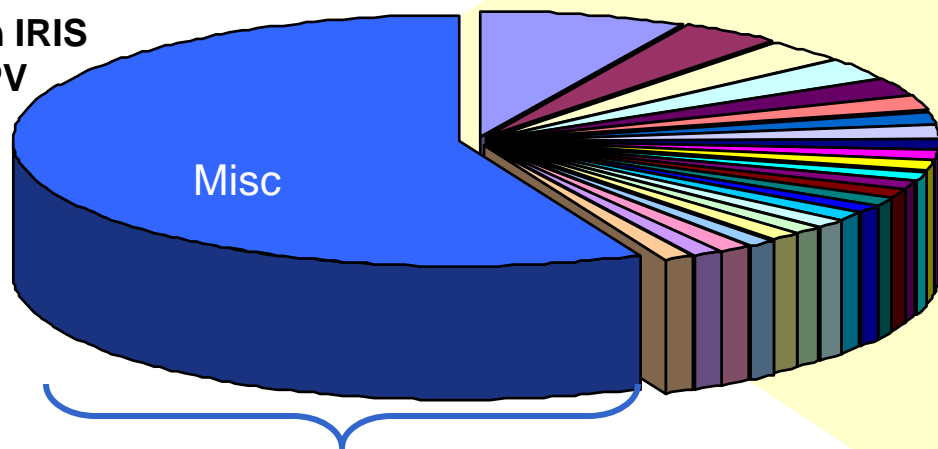
# The ToxCast Phase I Chemicals (320)

309 Unique Structures  
Replicates for QC

291 Pesticide Actives  
9 Industrial Chemicals  
8 Metabolites

56/73 Proposed Tier 1  
EDSP

122 in IRIS  
14 HPV



Classes with  
> 3 chemicals

Misc MOA classes with  
3 or fewer representatives

- Acetylcholine esterase inhibitors
- conazole fungicides
- Sodium channel modulators
- pyrethroid ester insecticides
- organothiophosphate acaricides
- dinitroaniline herbicides
- pyridine herbicides
- thiocarbamate herbicides
- imidazolinone herbicides
- organophosphate insecticides
- phenyl organothiophosphate insecticides
- aliphatic organothiophosphate insecticides
- amide herbicides
- aromatic fungicides
- chloroacetanilide herbicides
- chlorotriazine herbicides
- growth inhibitors
- organophosphate acaricides
- oxime carbamate insecticides
- phenylurea herbicides
- pyrethroid ester acaricides
- strobilurin fungicides
- unclassified acaricides
- unclassified herbicides

# ToxCast Assays

~500 Total Endpoints

## Cellular Assays

### Biochemical Assays

- Protein families
  - GPCR
  - NR
  - Kinase
  - Phosphatase
  - Protease
  - Other enzyme
  - Ion channel
  - Transporter
- Assay formats
  - Radioligand binding
  - Enzyme activity
  - Co-activator recruitment

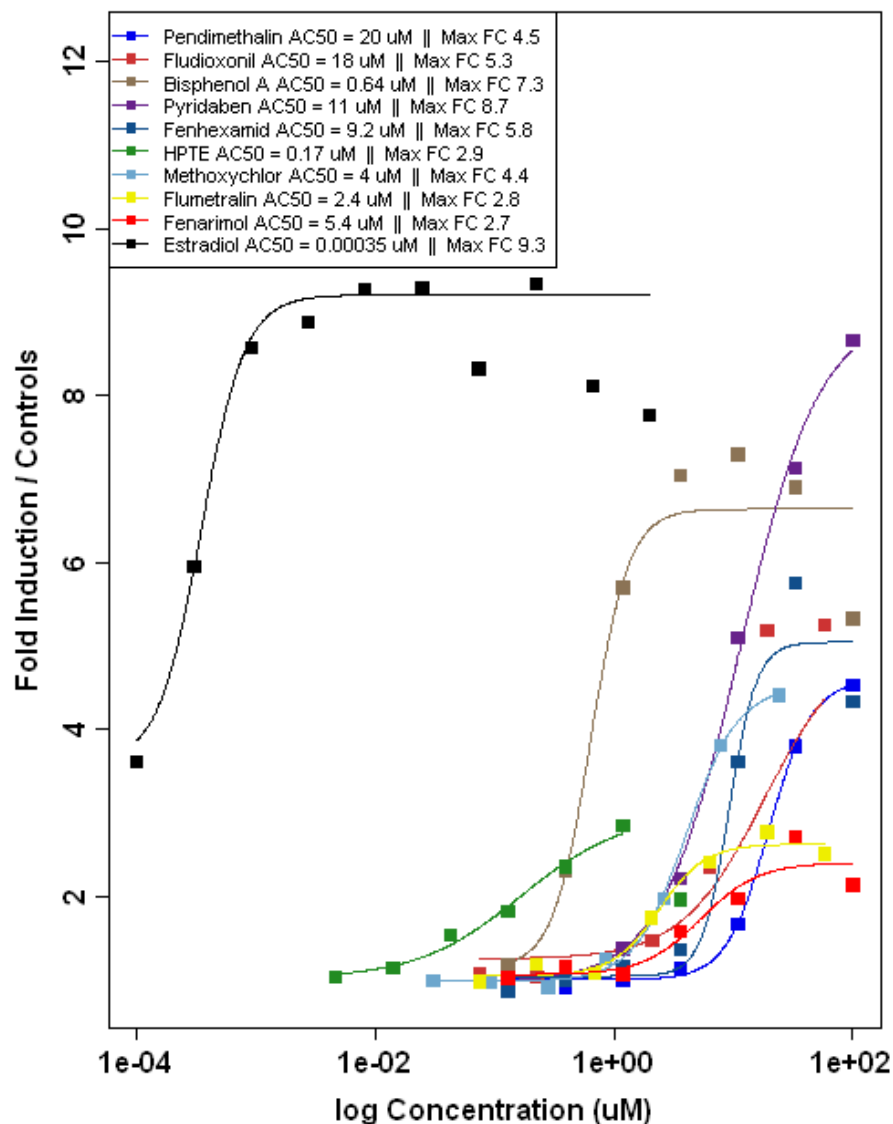
Primarily Human / Rat  
Exception: Zebrafish development (Stephanie Padilla)

- Cell lines
  - HepG2 human hepatoblastoma
  - A549 human lung carcinoma
  - HEK 293 human embryonic kidney
- Primary cells
  - Human endothelial cells
  - Human monocytes
  - Human keratinocytes
  - Human fibroblasts
  - Human proximal tubule kidney cells
  - Human small airway epithelial cells
  - Rat hepatocytes
  - Mouse embryonic stem cells (Sid Hunter)
- Biotransformation competent cells
  - Primary rat hepatocytes
  - Primary human hepatocytes
- Assay formats
  - Cytotoxicity
  - Reporter gene
  - Gene expression
  - Biomarker production
  - High-content imaging for cellular phenotype

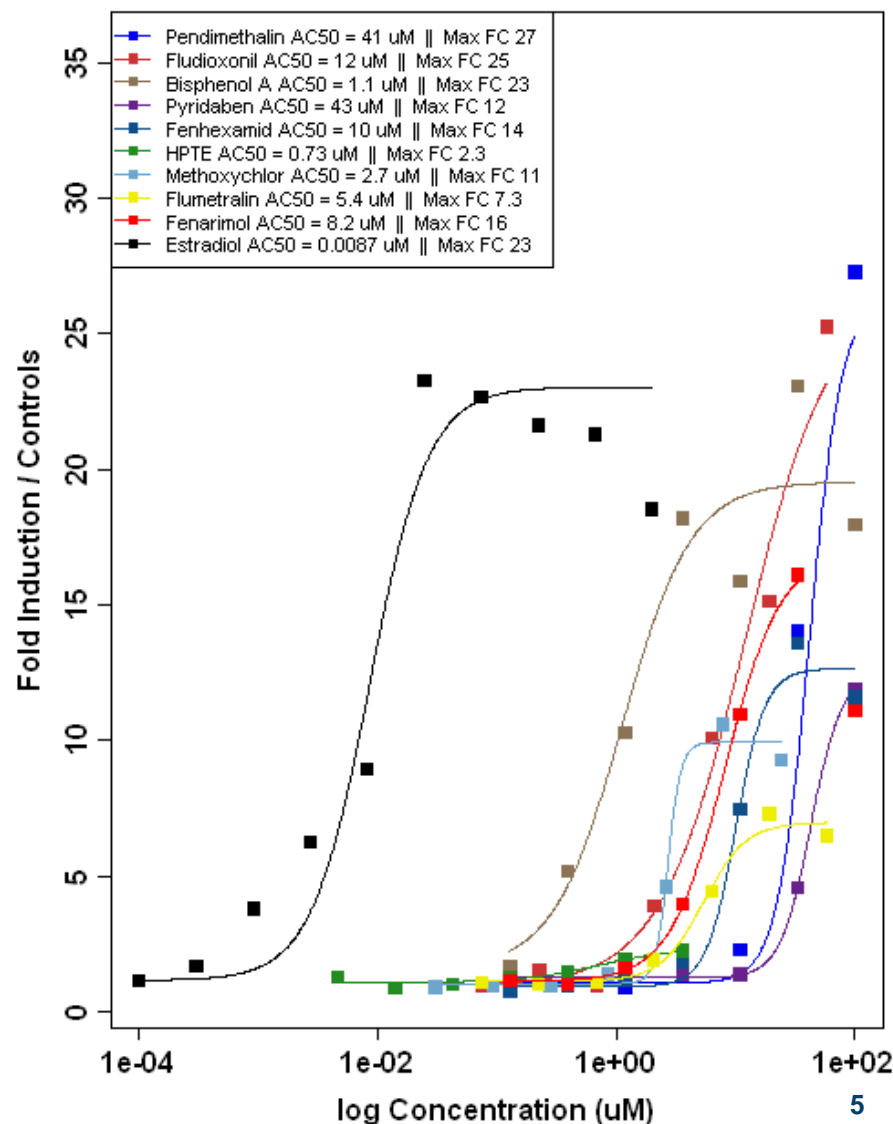
# Data Analysis:

## What is a hit?

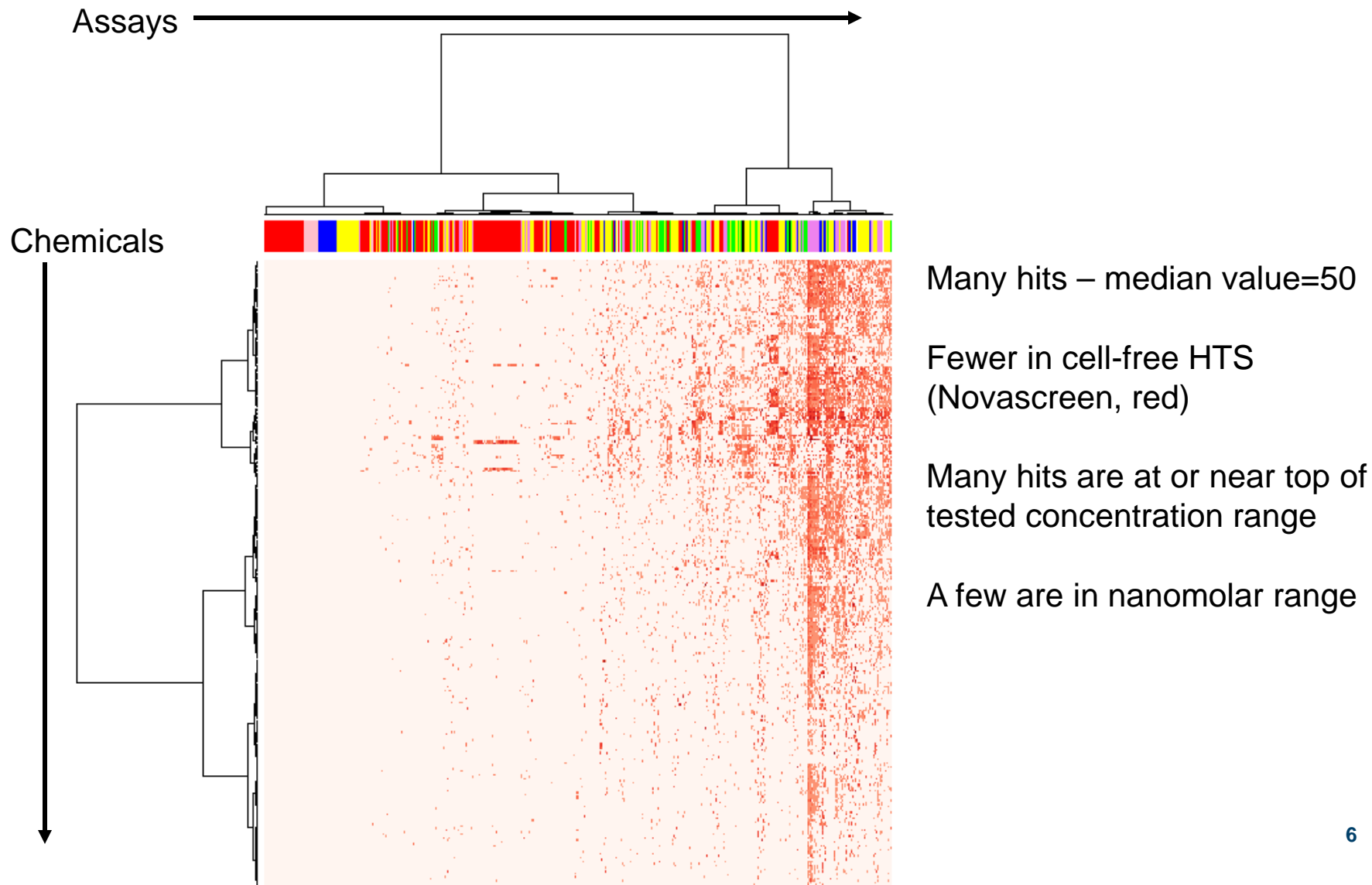
Attagene ERE\_CIS



Attagene ERa\_TRANS

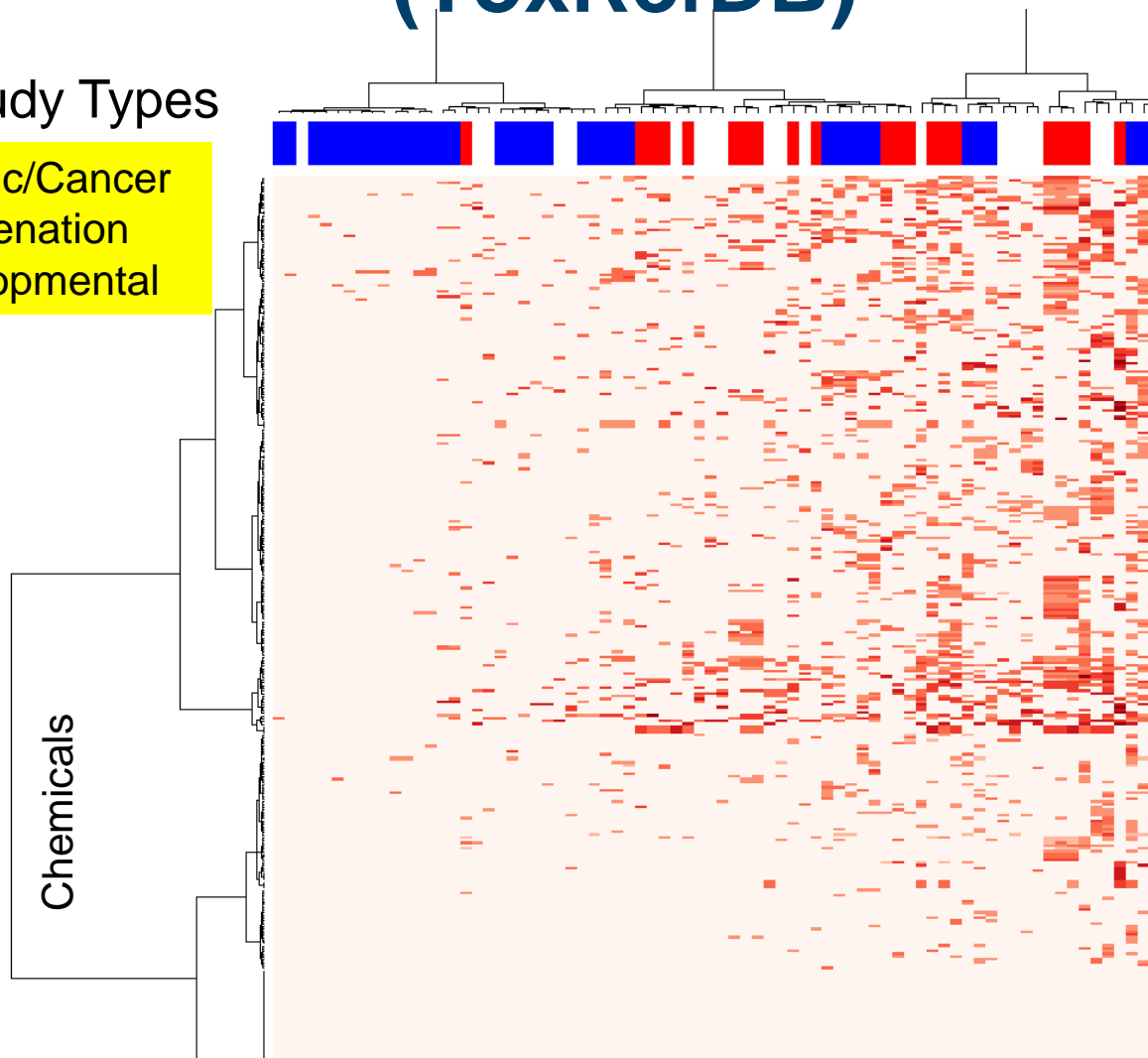
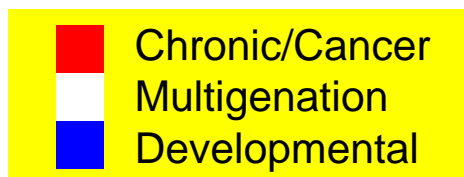


# The ToxCast In Vitro Data Set



# Reference Toxicity Database (ToxRefDB)

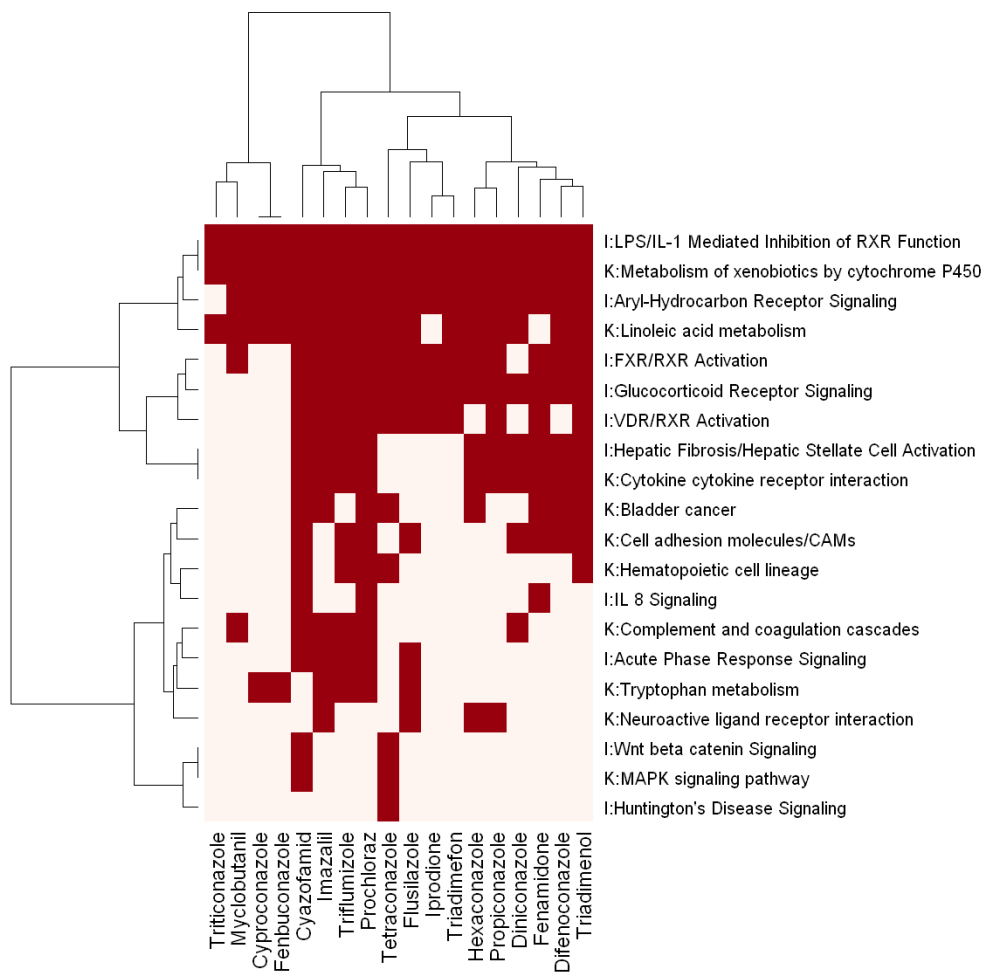
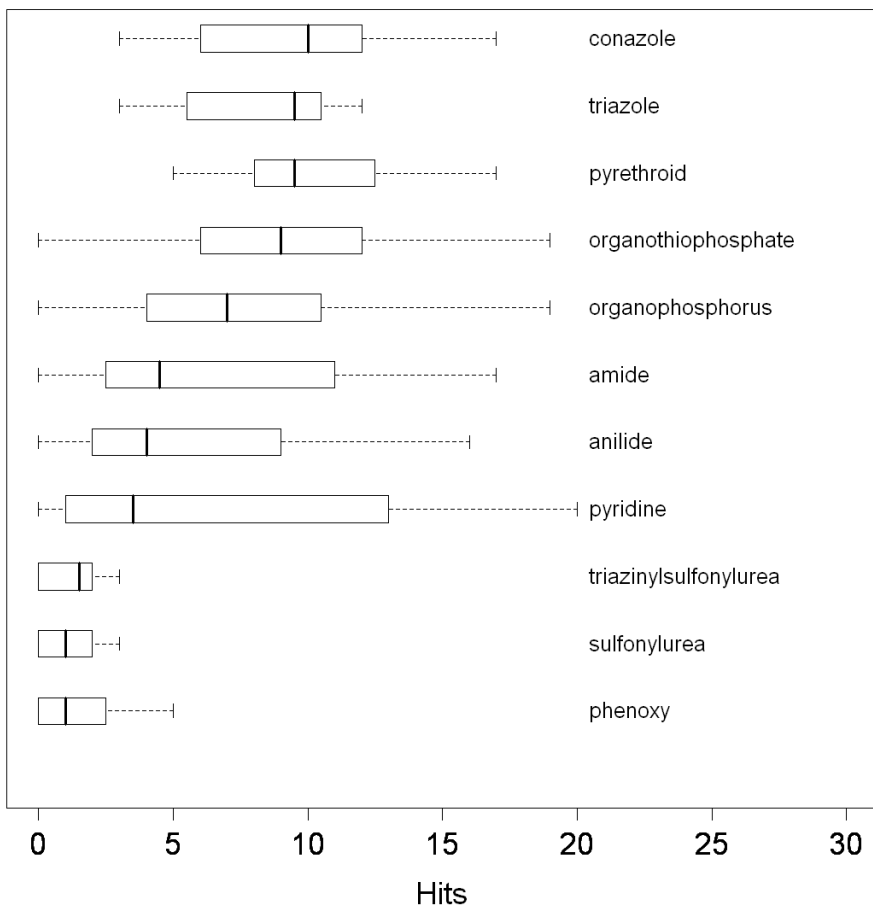
## In Vivo Study Types





# Chemicals show a wide range of activity

Minimal Pathways, 30 uM

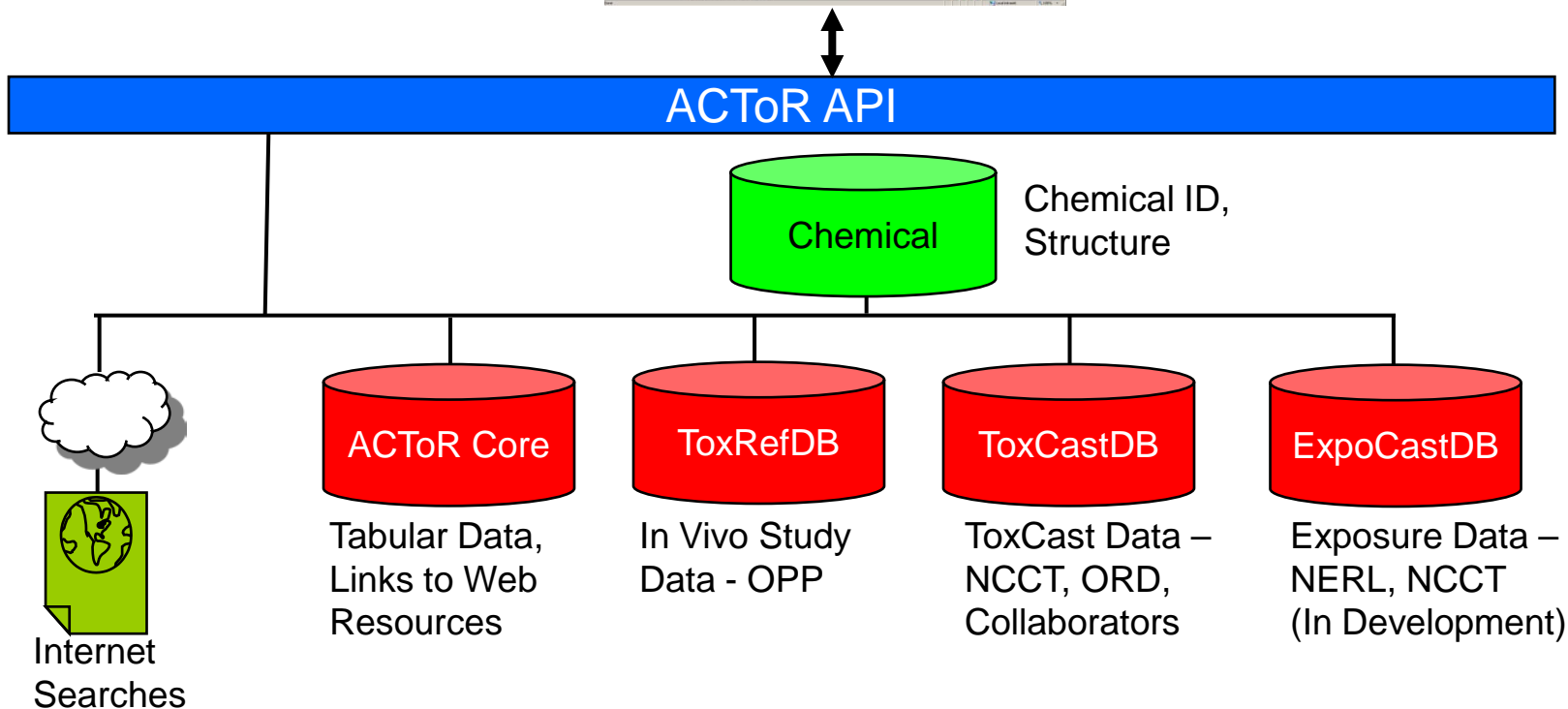
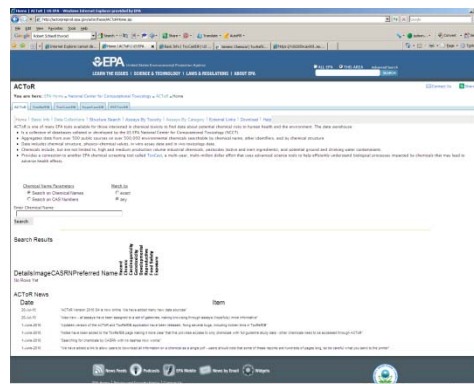


## Where is the data?

- ToxCastDB contains the ToxCast and ToxRefDB data for browsing
  - <http://actor.epa.gov/actor/faces/ToxMiner/Home.jsp>
- ACToR contains a much large amount of data on other types of toxicity and exposure
  - <http://actor.epa.gov>

# ACToR Aggregated Computational Toxicology Resource

<http://actor.epa.gov/>



# Data Definitions

- Substance
  - A chemical from one source
  - Name(s), CASRN
  - Source-specific unique ID
  - Assay Data
- Compound
  - Chemical structure from one source
  - Source-specific unique ID
- Generic Chemical
  - CASRN
  - Link to many substance (each with same CASRN)
  - Link to at most one compound
  - Links to all assay data from substances with same CASRN

# Data Definitions

- Assay
  - A collection of data on one or more substances
  - Comes from one data source
  - Can have several types of data included
  - Looks like an Excel spreadsheet
- Assay Component
  - One column of an assay table
- Assay Result
  - A data value for one substance and one assay component

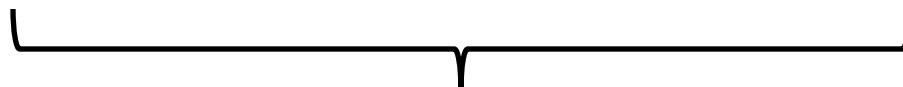
# Data Definitions

- Assay Phenotype
  - Type of disease associated with the assay
    - Carcinogenicity, GeneTox, ...
- Assay Category
  - Type of data: tabular, links to the web, human exposure
  - Allows assays to be grouped together
- Data Collection
  - A source of data
  - Substances
  - Compounds
  - Assays

## Data Sets

1. Attagene
2. ACEA
3. BioSeek
4. Cellumen
5. Gentronix
6. NCGC
7. Novascreen
8. CellzDirect
9. Solidus
10. ToxRefDB

- Perturbation Scores
  - Genes
  - Pathways
    - Ingenuity
    - KEGG
    - PathwayCommons
- PhysChem Properties
  - EPI Suite
  - LeadScope
  - QikProp
- Structure Classifiers




Not in current release





# Browsing ToxCast / Tox21 Data

<http://actor.epa.gov/actor/faces/ToxMiner/Home.jsp>


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**Chemical Name Parameters**  
☒ Enter Chemical Name:  
☐ Enter CAS Numbers:

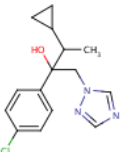
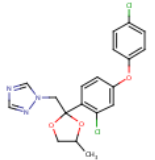
**Match by**  
☐ exact  
☒ any

Enter Chemical Name:

**Search**

---

### Chemical List

Details	Structure	Name	CASRN
<a href="#">Details</a>		Cyproconazole	94361-06-5
<a href="#">Details</a>		Difenoconazole	119446-68-3

Home page – search by  
name or CASRN

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**Chemical: Cyproconazole**

CASRN  
Smiles  
Source Name SID  
Source Name CID  
ACToR

94361-06-5  
CC1=CC=C(C(CN2C=NC=N2)(C(C)C3CC3)O)C=C1  
DSSTOX\_40409  
DSSTOX\_12601  
[Find in ACToR DB](#)

Source	Assay	Assay Name	Species	Gene	Value	Units
Attagene	<a href="#">ATG_ERa_TRANS</a>	Attagene Factorial trans Era	Homo sapiens	<a href="#">ESR1</a>	36.0	uM
Attagene	<a href="#">ATG_PXRE_CIS</a>	Attagene Factorial cis PXRE	Homo sapiens	<a href="#">NR1H2</a>	41.0	uM
BioSeek	<a href="#">BSK_BE3C_IP10_down</a>	BiEPI_IL1b_TNF_a_IFN_g_24_CXCL10_IP10_down	Homo sapiens	<a href="#">CXCL10</a>	40.0	uM
BioSeek	<a href="#">BSK_BE3C_TGFb1_down</a>	BiEPI_IL1b_TNF_a_IFN_g_24_TGF_beta1_down	Homo sapiens	<a href="#">TGFb1</a>	4.44	uM
BioSeek	<a href="#">BSK_hDFCGF_Proliferation_down</a>	HDFn_IL1b_TNF_a_IFN_g_EGF_FGF_PDGFbb_24_Proliferation_72hr_down	Homo sapiens		48.4	uM
BioSeek	<a href="#">BSK_SAg_PBMCCytotoxicity_down</a>	HUVEC_PBMCC_SEB_TSST_24_PBMCCytotoxicity_down	Homo sapiens		13.3	uM
BioSeek	<a href="#">BSK_SM3C_LDRLR_down</a>	SMC_IL1b_TNF_a_IFN_g_24_LDRLR_down	Homo sapiens	<a href="#">LDLR</a>	40.0	uM
BioSeek	<a href="#">BSK_LPS_PGE2_up</a>	HUVEC_PBMCC_LPS_24_PGE2_up	Homo sapiens	<a href="#">PTGER2</a>	40.0	uM
BioSeek	<a href="#">BSK_LPS_TNFa_up</a>	HUVEC_PBMCC_LPS_24_TNF_alpha_up	Homo sapiens	<a href="#">TNF</a>	13.3	uM
BioSeek	<a href="#">BSK_SAg_CD69_up</a>	HUVEC_PBMCC_SEB_TSST_24_CD69_up	Homo sapiens	<a href="#">CD69</a>	1.48	uM
Cellumen	<a href="#">CLM_MitoMembPot_1hr</a>	Cellumen Mito Mem Potential	Homo sapiens		31.6	uM
Cellumen	<a href="#">CLM_MitoMembPot_72hr</a>	Cellumen Mito Mem Potential	Homo sapiens		200.0	uM
CellzDirect	<a href="#">CLZD_CYP1A1_24</a>	CellzDirect CYP1A1	Homo sapiens	<a href="#">CYP1A1</a>	5.6	uM
CellzDirect	<a href="#">CLZD_CYP1A1_48</a>	CellzDirect CYP1A1	Homo sapiens	<a href="#">CYP1A1</a>	6.56	uM
CellzDirect	<a href="#">CLZD_CYP1A2_24</a>	CellzDirect CYP1A2	Homo sapiens	<a href="#">CYP1A2</a>	4.92	uM
CellzDirect	<a href="#">CLZD_CYP1A2_48</a>	CellzDirect CYP1A2	Homo sapiens	<a href="#">CYP1A2</a>	7.78	uM
CellzDirect	<a href="#">CLZD_CYP2B6_6</a>	CellzDirect CYP2B6	Homo sapiens	<a href="#">CYP2B6</a>	10.5	uM
CellzDirect	<a href="#">CLZD_CYP2B6_24</a>	CellzDirect CYP2B6	Homo sapiens	<a href="#">CYP2B6</a>	6.9	uM
CellzDirect	<a href="#">CLZD_CYP2B6_48</a>	CellzDirect CYP2B6	Homo sapiens	<a href="#">CYP2B6</a>	3.62	uM

Assay  
Set


AC50 and units

Assay  
Link

Assay  
Name

Gene  
Link to Entrez Gene

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## Assay: Attagene Factorial trans Era

<b>Assay Id:</b>	22
<b>Source</b>	Attagene
<b>Source Name AID</b>	ATG_Era_TRANS
<b>Name</b>	Attagene Factorial trans Era
<b>Description</b>	Factorial reporter gene assay
<b>Number of Substances</b>	320
<b>Number of Components</b>	1
<b>Species</b>	Homo sapiens

Parameter	Value
ASSAY URL	<a href="#">Link Out</a> <a href="#">EXIT Disclaimer</a>
ASSAY CATEGORY	In vitro (Cellular)
ASSAY TARGET	ERa
ASSAY TARGET FAMILY	Transcription Factor
ASSAY TARGET SOURCE	Cell line
ASSAY TARGET SOURCE TYPE	HepG2
ASSAY GENE ID	2099
ASSAY GENE NAME	<a href="#">ESR1</a>
ASSAY TECHNOLOGY	Reporter gene assay
ASSAY MODE	DNA sequencer
ASSAY REFERENCE COMPOUND	"17b-Estradiol, Diethylstilbestrol(DES)"
ASSAY NOTE	Multiplexed reporter gene assay; Nuclear receptor pathway


  

Data		
Name	CASRN	ATG_Era_TRANS (uM)
HPTE	2971-36-0	0.73
2-Phenylphenol	90-43-7	43.0
Azinphos-methyl	86-50-0	47.0
Azoxystrobin	131860-33-8	17.0
Benfluralin	1861-40-1	36.0
Bensulide	741-58-2	2.0
Bifenthrin	82657-04-3	37.0
Bifenox A	80-05-7	1.1

Description of assay

AC50s for all positive chemicals

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### Data Collection: NCGC

Name: NCGC  
Description: NCGC - nuclear receptor assays  
Number of Chemicals: 320  
Number of Assays: 19  
Number of Data Points: 6080

#### Data Collection Summary


Name	Assay Name	Description	Substances	Components	Species	Gene
<a href="#">NCGC_AR_Agonist</a>	NCGC Reporter Gene Assay AR Agonist	GAL4 BLAM Reporter gene assay: AR	320	1	Homo sapiens	<a href="#">AR</a>
<a href="#">NCGC_AR_Antagonist</a>	NCGC Reporter Gene Assay AR Antagonist	GAL4 BLAM Reporter gene assay: AR	320	1	Homo sapiens	<a href="#">AR</a>
<a href="#">NCGC_ERalpha_Agonist</a>	NCGC Reporter Gene Assay ERa Agonist	GAL4 BLAM Reporter gene assay: ERa	320	1	Homo sapiens	<a href="#">ESR1</a>
<a href="#">NCGC_ERalpha_Antagonist</a>	NCGC Reporter Gene Assay ERa Antagonist	GAL4 BLAM Reporter gene assay: ERa	320	1	Homo sapiens	<a href="#">ESR1</a>
<a href="#">NCGC_FXR_Agonist</a>	NCGC Reporter Gene Assay FXR Agonist	GAL4 BLAM Reporter gene assay: FXR	320	1	Homo sapiens	<a href="#">NR1H4</a>
<a href="#">NCGC_GR_Agonist</a>	NCGC Reporter Gene Assay GR Agonist	GAL4 BLAM Reporter gene assay: GR	320	1	Homo sapiens	<a href="#">NR3C1</a>
<a href="#">NCGC_HEK293_Viability</a>	NCGC_HEK293_Viability	NCGC_HEK293_Viability	320	1	Homo sapiens	
<a href="#">NCGC_LXR_Agonist</a>	NCGC Reporter Gene Assay LXRb Agonist	GAL4 BLAM Reporter gene assay: LXRa	320	1	Homo sapiens	<a href="#">NR1H2</a>
<a href="#">NCGC_p53</a>	NCGC p53 assay	NCGC p53 assay	320	1	Homo sapiens	<a href="#">TP53</a>
<a href="#">NCGC_PPARa_Agonist</a>	NCGC Reporter Gene Assay PPARa Agonist	GAL4 BLAM Reporter gene assay: PPARa	320	1	Homo sapiens	<a href="#">PPARA</a>
<a href="#">NCGC_PPARd_Agonist</a>	NCGC Reporter Gene Assay PPARd Agonist	GAL4 BLAM Reporter gene assay: PPARd	320	1	Homo sapiens	<a href="#">PPARD</a>
<a href="#">NCGC_PPARg_Agonist</a>	NCGC Reporter Gene Assay PPARg Agonist	GAL4 BLAM Reporter gene assay: PPARg	320	1	Homo sapiens	<a href="#">PPARG</a>
<a href="#">NCGC_PXR_Agonist_human</a>	NCGC Reporter Gene Assay PXR Agonist (Human)	GAL4 BLAM Reporter gene assay: PXR Human	320	1	Homo sapiens	<a href="#">NR1I2</a>
<a href="#">NCGC_PXR_Agonist_rat</a>	NCGC Reporter Gene Assay PXR Agonist (Rat)	GAL4 BLAM Reporter gene assay: PXR Rat	320	1	Rattus norvegicus	<a href="#">Nr1i2</a>
<a href="#">NCGC_RXRa_Agonist</a>	NCGC Reporter Gene Assay RXRa Agonist	GAL4 BLAM Reporter gene assay: RXRa	320	1	Homo sapiens	<a href="#">RXRA</a>
<a href="#">NCGC_TRbeta_Agonist</a>	NCGC Reporter Gene Assay TRb Agonist	GAL4 BLAM Reporter gene assay: TRb	320	1	Homo sapiens	<a href="#">THRB</a>
<a href="#">NCGC_TRbeta_Antagonist</a>	NCGC Reporter Gene Assay TRb Antagonist	GAL4 BLAM Reporter gene assay: TRb	320	1	Homo sapiens	<a href="#">THRB</a>
<a href="#">NCGC_VDR_Agonist</a>	NCGC Reporter Gene Assay VDR Agonist	GAL4 BLAM Reporter gene assay: VDR	320	1	Homo sapiens	<a href="#">VDR</a>
<a href="#">NCGC_AhR</a>	NCGC Reporter Gene Assay AHR Agonist	GAL4 BLAM Reporter gene assay: AHR	320	1	Homo sapiens	<a href="#">AHR</a>

List of all assays from a particular data set

# Genes associated with assays

Lists all assays associated with each gene

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Busy Sorting Data:

Genes With Assays			
Name	Symbol	Species	Assays
aryl hydrocarbon receptor	AHR	Homo sapiens	ATG_Ahr_CIS NCGC_AhR
jun oncogene	JUN	Homo sapiens	ATG_AP_1_CIS StressKinase
transcription factor AP-2 alpha (activating enhancer binding protein 2 alpha)	TFAP2A	Homo sapiens	ATG_AP_2_CIS
androgen receptor	AR	Homo sapiens	ATG_AR_TRANS NCGC_AR_Agonist NCGC_AR_Antagonist NVS_NR_hAR
bone morphogenetic protein receptor, type II (serine/threonine kinase)	BMPR2	Homo sapiens	ATG_BRE_CIS
CCAAT/enhancer binding protein (C/EBP), beta	CEBPB	Homo sapiens	ATG_C_EBP_CIS
nuclear receptor subfamily 1, group I, member 3	NR1I3	Homo sapiens	ATG_CAR_TRANS ATG_PBREM_CIS NVS_NR_hCAR_Antagonist
cAMP responsive element binding protein 3	CREB3	Homo sapiens	ATG_CRE_CIS
nuclear receptor subfamily 1, group H, member 3	NR1H3	Homo sapiens	ATG_DR4_LXR_CIS ATG_LXRa_TRANS
tumor necrosis factor receptor superfamily, member 10b	TNFRSF10B	Homo sapiens	ATG_DR5_CIS
upstream transcription factor 1	USF1	Homo sapiens	ATG_E_Box_CIS
E2F transcription factor 1	E2F1	Homo sapiens	ATG_E2F_CIS
early growth response 1	EGR1	Homo sapiens	ATG_EGR_CIS
estrogen receptor 1	ESR1	Homo sapiens	ATG_ERa_TRANS ATG_ERE_CIS NCGC_ERalpha_Agonist NCGC_ERalpha_Antagonist NVS_NR_hER

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## Coming Soon

- Mouse Embryonic Stem Cell Data (With Sid Hunter, EPA NHEERL)
- RTK Data – Reverse Toxicokinetics (with R. Thomas, Hamner)
- ExpoCastDB (With collaborators in EPA NERL)





- Data Sets: Bio-centric View
  - Assay overview
  - How were the data calculated?
  - How did the replicates perform?
  - Example results
  - What did the assay measure?
  - Potential assay artifacts

# ToxCast Phase I Data Sets

- Biochemical
  - Novascreen
- Cell-Based
  - ACEA
  - Attagene
  - BioSeek
  - Cellumen
  - CellzDirect
  - Gentronix
  - NCGC
  - Solidus

# Novascreen: 239 Biochemical Assays (Abstract 59)

- Protein super-families
  - GPCR
  - Kinase
  - Phosphatase
  - Protease
  - Ion channel
  - Nuclear receptor
  - Other enzyme
  - CYP P450 inhibition
- Various formats:
  - Radioligand receptor binding
  - Fluorescent receptor binding
  - Fluorescent enzyme substrate-intensity quench
  - Fluorescent enzyme substrate-mobility shift
- Initial screening:
  - 25  $\mu$ M in duplicate
  - 10  $\mu$ M in duplicate (CYPs)
- Normalize data to assay window
  - % of control activity (central reference – scalar reference)

# Novascreen: What do the assays measure?

- Mainly direct effects of chemical on target protein
  - Enzyme activity
  - Ligand binding
- False positives:
  - Fluorescent compounds—fluorescing and quenching
  - Reactive compounds/covalent modification of target
  - Physical effects—colloid aggregation of target
  - Operational
- False negatives:
  - Solubility
  - Inappropriate assay conditions
  - Operational
  - Target protein not physiological
  - Lack of biotransformation

# Data Correction Algorithm Examples (Additive)

AChE

	1	2	3	4	5	6	7	8	9	10	11	12
A	-3	3	-29	-100	-98	-100	-68	-68	-89	-89	-98	-98
B	-4	7	-29	-12	-10	-12	-27	-22	-30	-30	-25	-16
C	8	2	-23	-23	-19	-26	-25	-26	-26	-28	-25	-13
D	-14	1	-17	-21	-23	-25	-18	-26	-26	-27	-25	-17
E	-11	-5	-23	-25	-16	-20	-26	-22	-27	-30	-26	-13
F	-16	0	-7	-7	-17	-21	-22	-24	-29	-33	-19	-17
G	-5	4	-32	-18	-25	-17	-25	-28	-25	-32	-20	-13
H	-85	-78	-22	-22	2	-14	-17	-17	NaN	NaN	NaN	NaN

normalized

	1	2	3	4	5	6	7	8	9	10	11	12
A	-1	1	-19	-100	-98	-100	-67	-67	-89	-89	-98	-98
B	0	7	-20	-10	-2	0	2	9	-15	-16	-9	-4
C	10	10	-3	-3	0	-6	-2	-5	-16	-18	-11	-1
D	4	14	7	2	2	-5	6	0	-13	-11	-9	-3
E	7	13	4	0	7	4	-2	2	-15	-18	-10	-1
F	5	22	16	16	9	6	3	3	-15	-17	-3	-2
G	12	23	-4	6	1	5	3	0	-11	-16	-1	4
H	-52	-44	7	9	20	8	4	5	NaN	NaN	NaN	NaN

Normalized  
& corrected

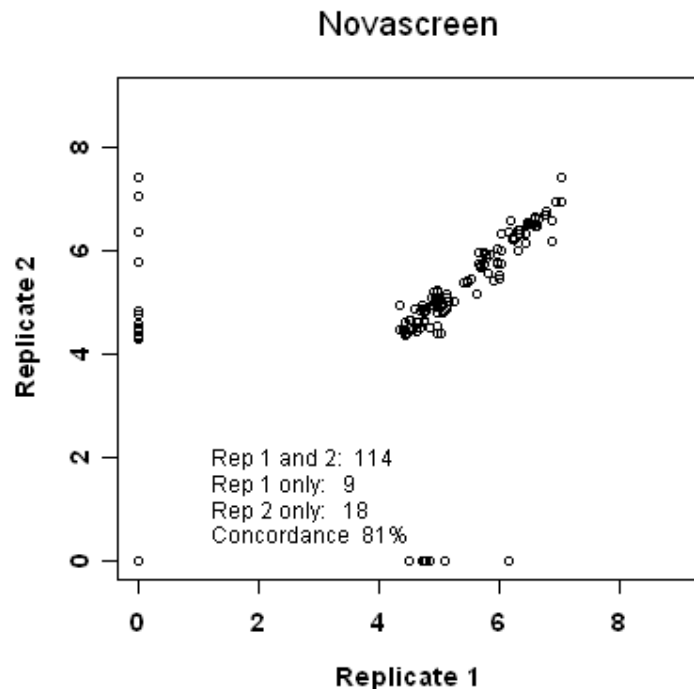
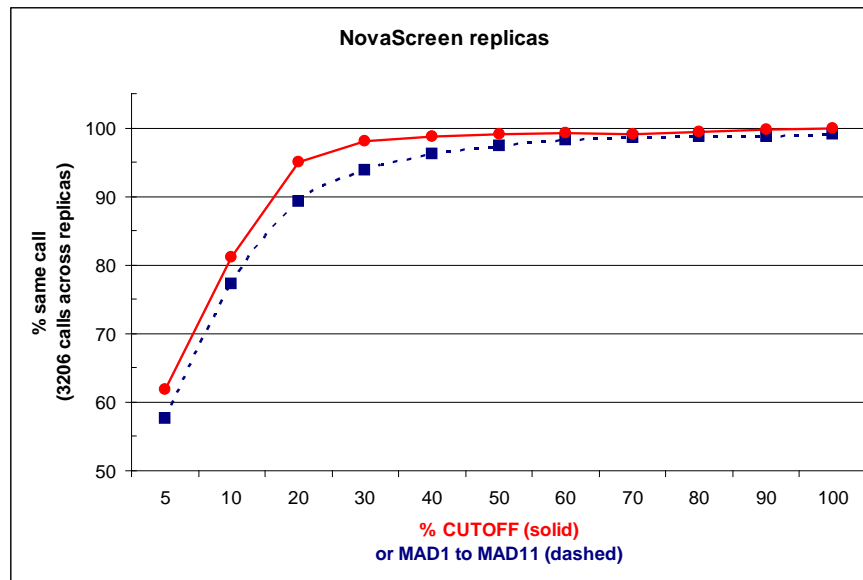
Caspase 10

	1	2	3	4	5	6	7	8	9	10	11	12
A	25	0	-3	-97	-100	-101	-35	-33	-76	-80	-100	-99
B	9	-17	-16	-23	-28	-17	-13	-16	-22	-15	-22	-3
C	3	-22	-10	-27	-24	-20	-39	-31	-24	-28	-26	-6
D	15	-22	-15	-27	-23	-27	-13	-15	-17	-18	-20	-3
E	3	-16	-16	-17	-18	-7	-6	-13	2	5	-17	12
F	-11	-34	-18	-26	-29	-21	-22	-19	-18	-19	-25	0
G	-23	-30	-27	-24	-17	-6	-8	-11	-22	-12	-9	10
H	-15	-28	-17	-20	-14	-28	-5	4	-19	-4	-18	5

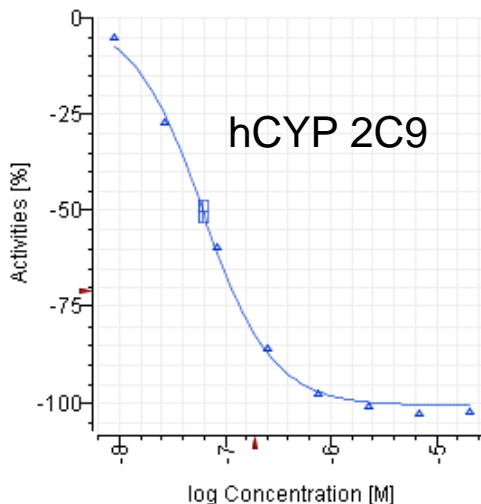
	1	2	3	4	5	6	7	8	9	10	11	12
A	23	0	-3	-97	-100	-101	-36	-34	-76	-81	-100	-99
B	13	6	7	0	-6	-3	3	2	-6	1	-5	0
C	6	-1	10	-7	-4	-5	-21	-13	-5	-10	-7	-5
D	18	0	2	-7	0	-7	6	4	0	-1	-4	1
E	1	1	2	-3	-5	-2	7	0	14	17	-6	9
F	-9	-12	4	-3	-11	-9	-4	-1	-7	-4	-11	-1
G	-17	-5	-5	-4	-1	-1	13	6	-2	3	5	8
H	-9	-11	2	0	4	-14	9	15	-4	10	-7	8

# Novascreen Concentration-Response Testing

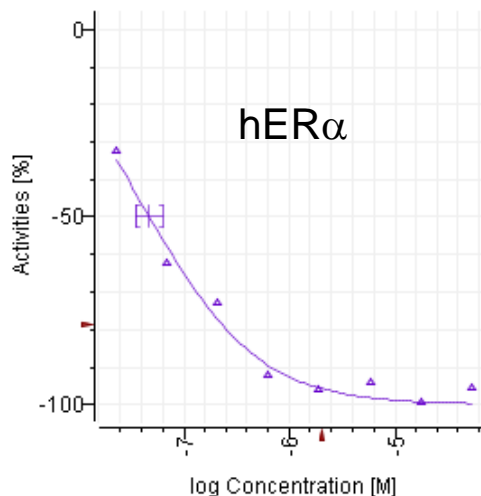
- Retest actives:
  - Median absolute deviation (MAD)  
median  $1x-x_{med}$   
two MADs or 30% activity
  - 8 conc/3-fold serial dilutions
    - 50  $\mu$ M high conc
    - 25  $\mu$ M high conc for CYPs
- Normalize to assay window
- Fit % Activity data to 3- or 4-parameter Hill function
  - Sometimes had to fix top or bottom of curve
  - Did not extrapolate beyond testing range
  - Manual or automated removal of obvious outliers



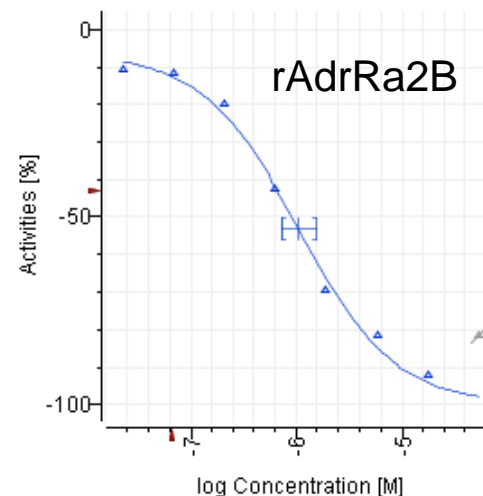
# Novascreen: Example Curve Fits



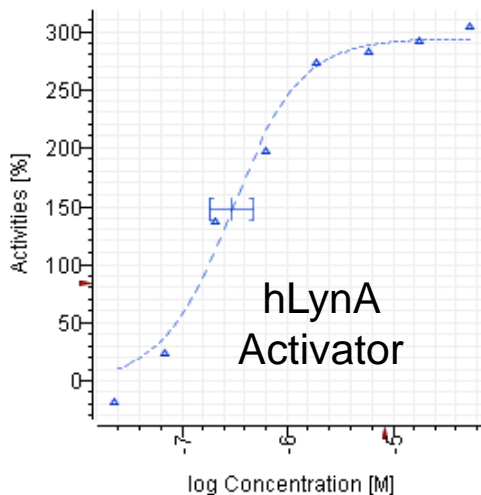
■ TV000541: Cyproconazole



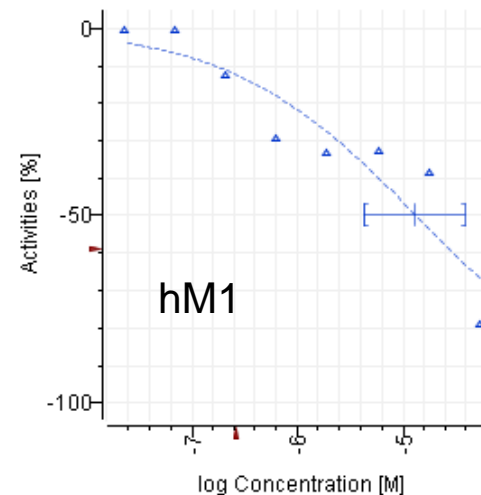
■ TV000626: 2,2-Bis(4-hydroxyphenyl)-1,1,1-trichloroethane



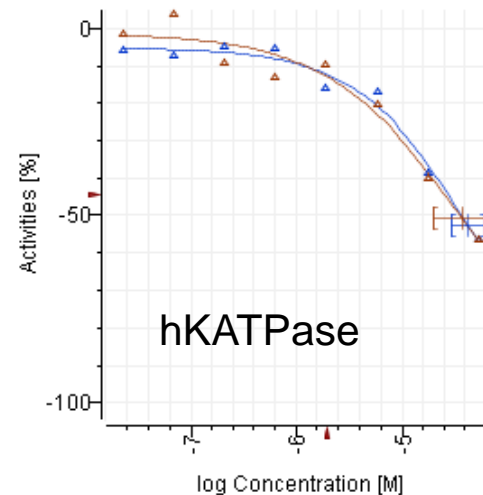
■ TV000551: Amitraz



■ TV000783: Mancozeb



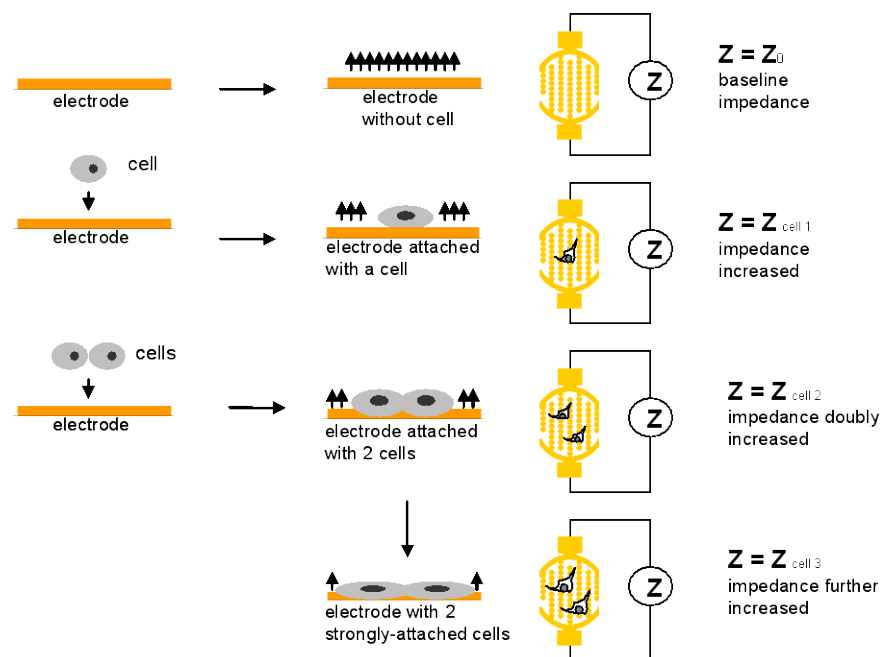
■ TV000526: Paclitaxel



■ TV000517: 3-Iodo-2-propynylbutyl carbamate  
■ TV000719: 3-Iodo-2-propynylbutyl carbamate

# ACEA: Real Time Cell Growth Kinetics

- Cytotoxicity with potential mechanistic interpretation
- Human A549 lung carcinoma cell line
  - ACEA experience with line
  - Reference compound effects
- Concentration-response testing
  - 8 conc/3-fold serial dilutions
  - Duplicate wells
- Real-time measurements during exposure (0-72 hr)
- IC50 and LELs calculated



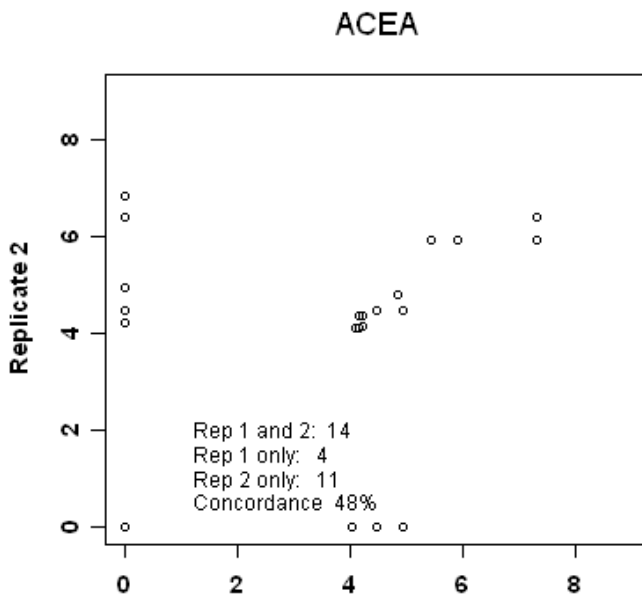


# ACEA: What is measured?

- General cytotoxicity in transformed cell line
- False positives
  - No obvious
  - Operational
- False negatives
  - Operational error
  - Solubility
  - Lack of appropriate toxicity targets
  - Lack of biotransformation

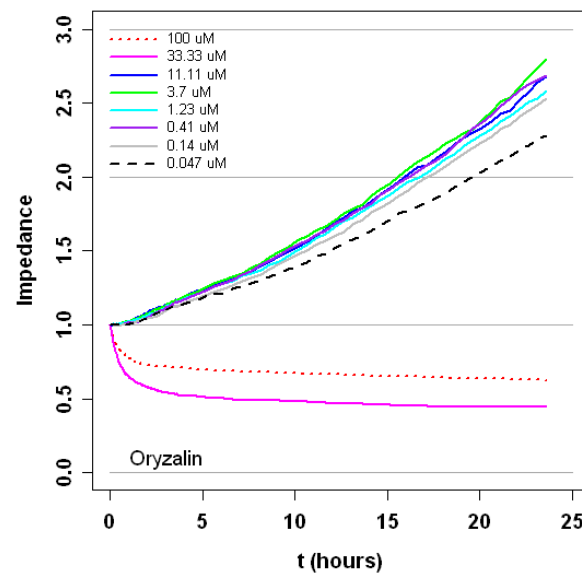
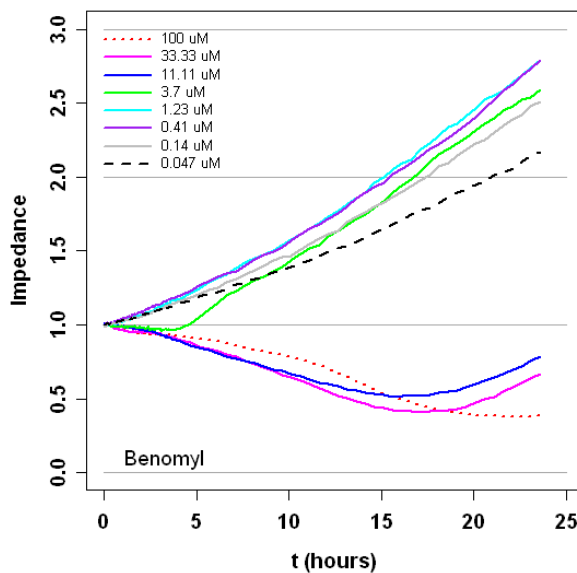
# ACEA: Data examples

## Replicate Analysis:



TV000019

TV000079



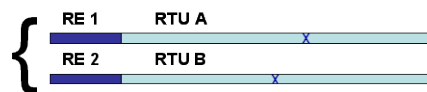
## Example Plots:

# Attagene Multiplexed Transcription Factor Assays

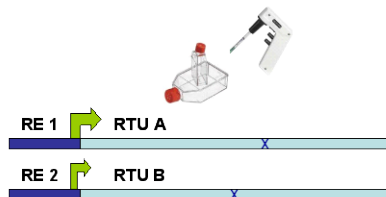
- Modulation of TF activity in human hepatoma HepG2 cells
- Multiplexed reporter gene assay
  - *cis* 52 assays (response element driving reporter)
  - *trans* 29 assays (GAL4-NR\_LBD driving reporter) “ligand detection”
- IC50 for cytotoxicity measured first in HepG2
- High concentration either 100  $\mu$ M or 1/3 calculated IC50 for cytotoxicity
- Seven concentrations, 3-fold serial dilutions, 24 hr exposure
- Cells harvested, RNA isolated, processed for reporter gene quantitation
- LEL provided in data set

## Multiplexed Reporter Gene Assay

Library of RTUs



Cell Transfection



Transcription

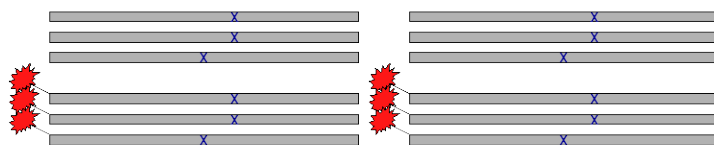
RNA Isolation



Reverse transcription



PCR amplification

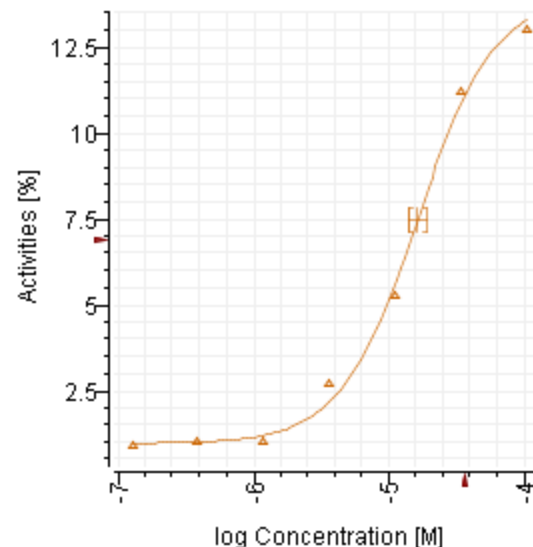
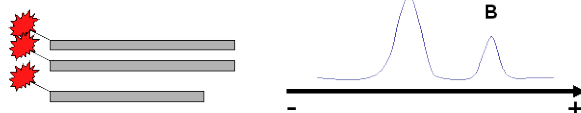


Labeling

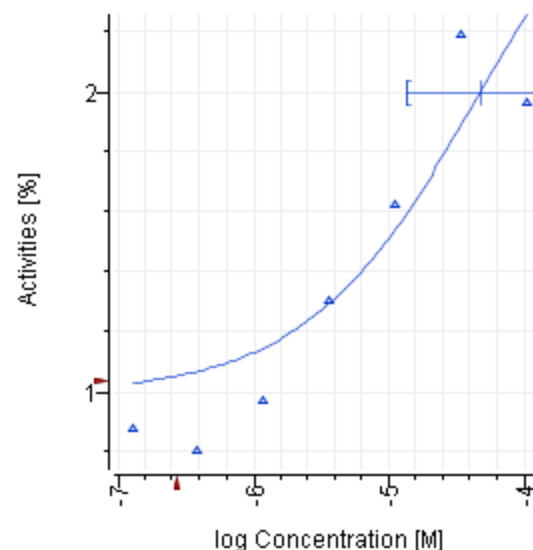
Processing (Hpa I)



Separation and detection  
(capillary electrophoresis)



TV000392

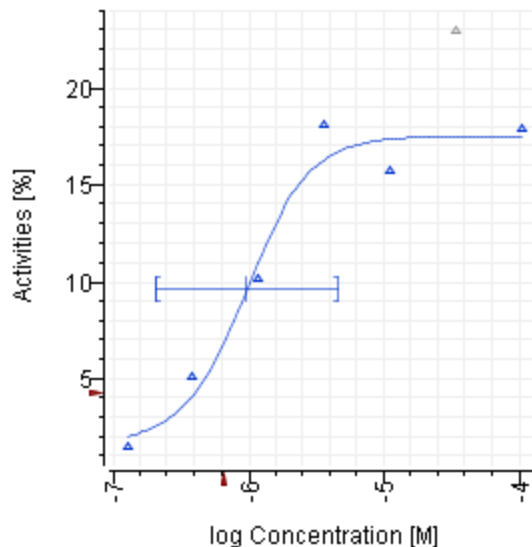


# Attagene: What Is Being Measured?

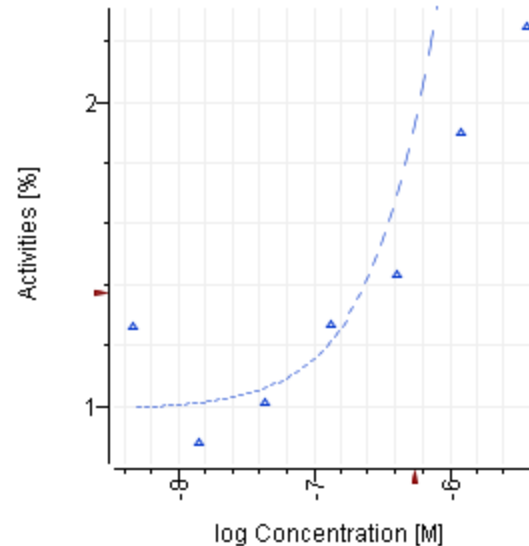
- cis Assays
  - Up/down regulation of endogenous transcription factor activity in transformed cell line
  - False positives
    - General cytotoxic response resulting in non-specific transcriptional activity
    - Promiscuity of response elements
    - Statistical, not biologically, significant response
    - Operational
  - False negatives
    - Solubility
    - Cytotoxicity
    - Operational
    - Lack of endogenous machinery
    - Lack of biotransformation
- trans Assays
  - NR agonist activity
  - False positives
    - General cytotoxic response resulting in non-specific transcriptional activity
    - Statistical, not biologically, significant response
    - Operational
  - False negatives
    - Solubility
    - Cytotoxicity
    - Operational
    - Lack of endogenous machinery
    - Lack of biotransformation

# Attagene: corresponding *cis* and *trans* assays

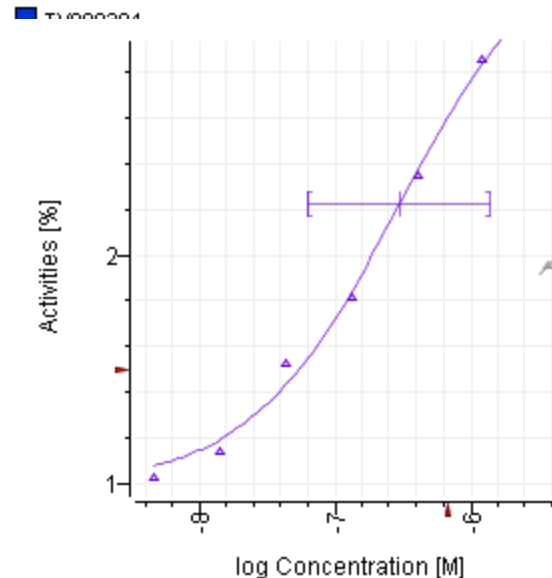
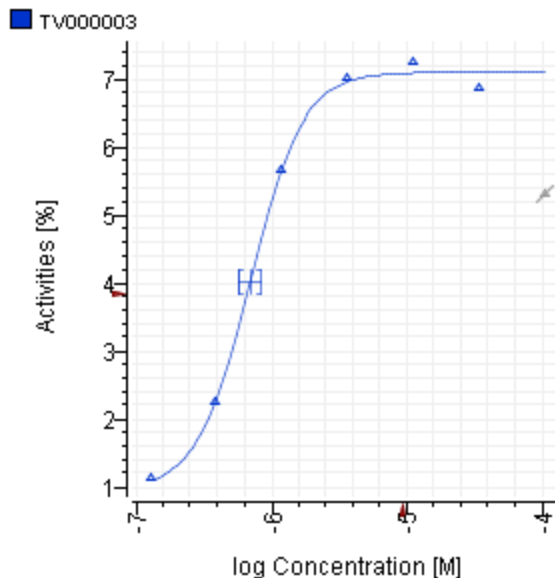
**Bisphenol A**



**HPTE**



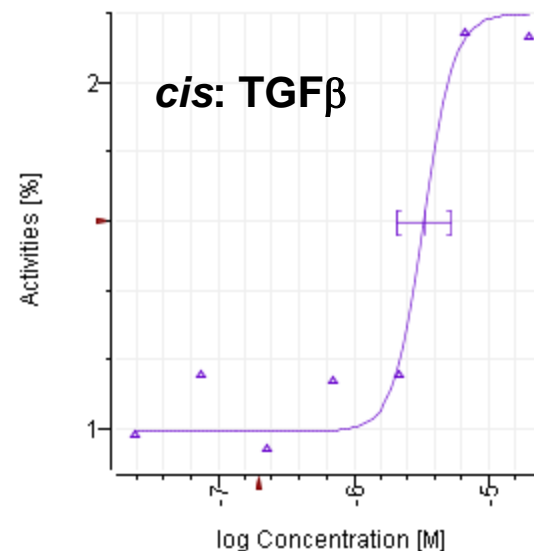
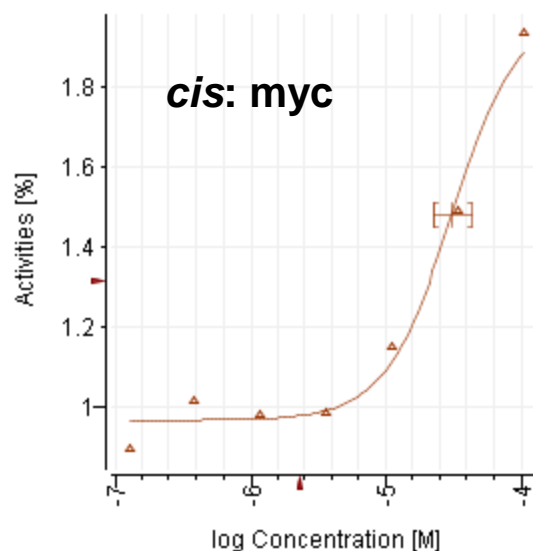
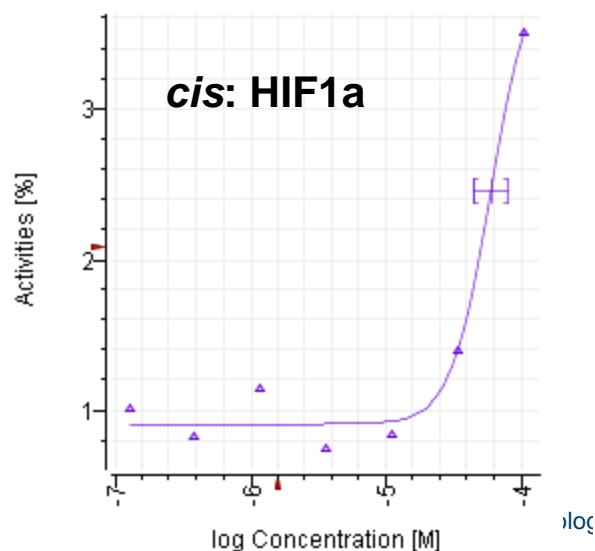
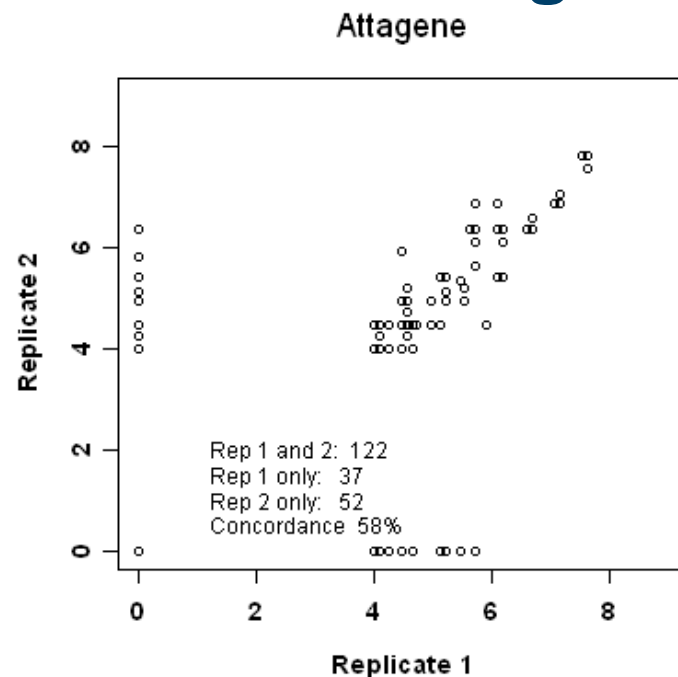
***trans*: ERa**



***cis*: ERE**

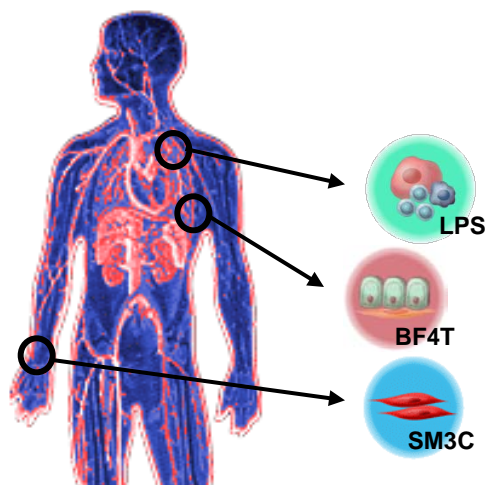
# Attagene: Data calculation challenges

- No positive reference compound for each endpoint
- Responses, especially for *cis* assays, tended to be monotonic so potency value difficult to derive
- Biological vs statistical significance for LEL



# BioSeek: BioMAP® Technology Platform (Abstract 24)

## Assays



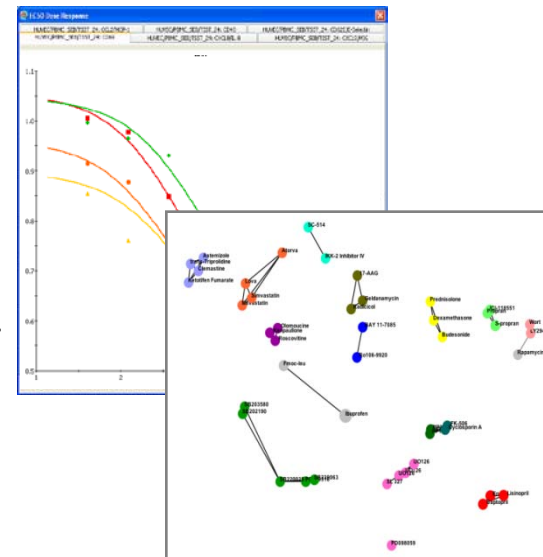
Human primary cells  
Disease-like culture  
conditions

## Profile Database



Biological responses to  
drugs and stored in the  
database

## Informatics








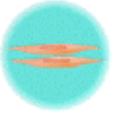


Specialized informatics tools  
are used to mine and analyze  
biological data

***Primary Human Cell-Based Assay Platform for Human Pharmacology***





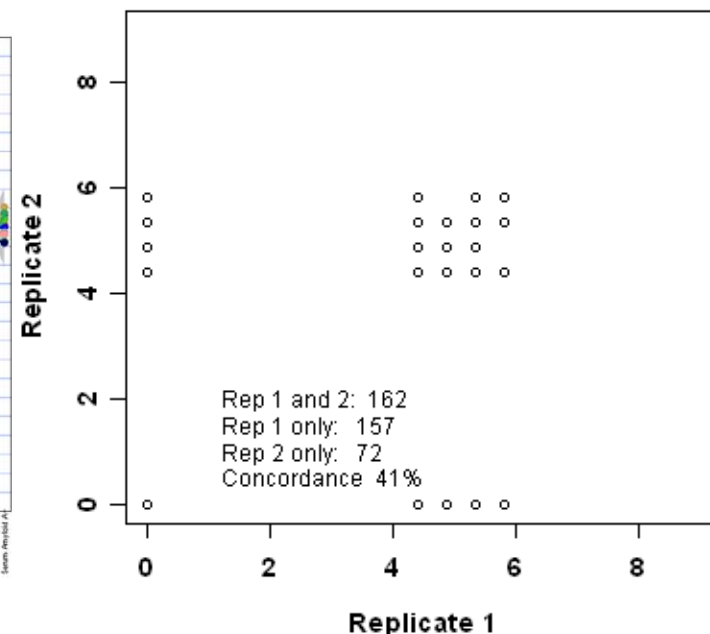
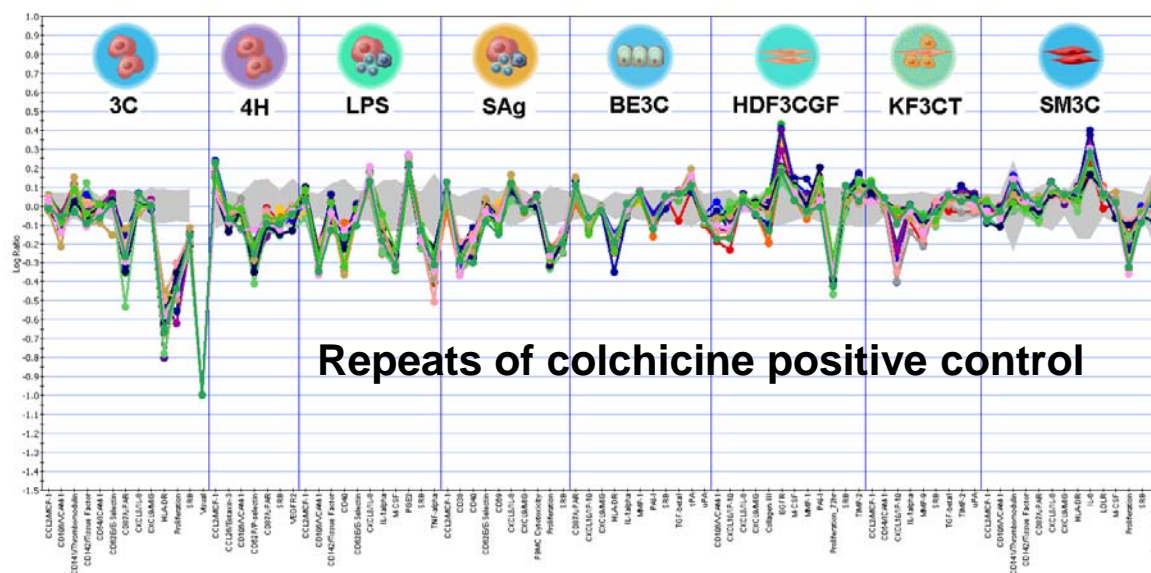
# BioSeek Assays Tested Against ToxCast\_320

System		Cell Types	Environment	Readouts
3C		Endothelial cells	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$	MCP-1, VCAM-1, ICAM-1, Thrombomodulin, Tissue Factor, E-selectin, uPAR, IL-8, MIG, HLA-DR, Prolif., Vis., SRB (13)
4H		Endothelial cells	IL-4+histamine	VEGFRII, P-selectin, VCAM-1, uPAR, Eotaxin-3, MCP-1, SRB (7)
LPS		Peripheral Blood Mononuclear Cells + Endothelial cells	TLR4	CD40, VCAM-1, Tissue Factor, MCP-1, E-selectin, IL-1a, IL-8, M-CSF, TNF-a, PGE2, SRB (11)
SAg		Peripheral Blood Mononuclear Cells + Endothelial cells	TCR	MCP-1, CD38, CD40, CD69, E-selectin, IL-8, MIG, PBMC Cytotox., SRB, Proliferation (10)
BE3C		Bronchial epithelial cells	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$	uPAR, IP-10, MIG, HLA-DR, IL-1a, MMP-1, PAI-1, SRB, TGF-b1, tPA, uPA (11)
HDF3CGF		Fibroblasts	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$ +bFGF+EGF+PDGF-BB	VCAM-1, IP-10, IL-8, MIG, Collagen III, M-CSF, MMP-1, PAI-1, Proliferation, TIMP-1, EGFR, SRB (12)
KF3CT		Keratinocytes + Fibroblasts	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$ +TGF- $\beta$	MCP-1, ICAM-1, IP-10, IL-1a, MMP-9, TGF-b1, TIMP-2, uPA, SRB (9)
SM3C		Vascular smooth muscle cells	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$	MCP-1, VCAM-1, Thrombomodulin, Tissue Factor, IL-6, LDLR, SAA, uPAR, IL-8, MIG, HLA-DR, M-CSF, Prolif., SRB (14)

# BioSeek Assays

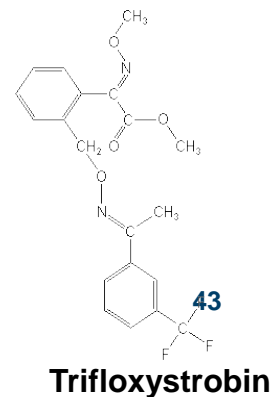
- Chemicals tested at 4 concentrations: 40, 13.3, 4.4, 1.5  $\mu$ M, single well
- Exposure started 1 hr before stimulation of cell signaling pathways
- Following 24 hr exposure, endpoints measured by ELISA (also Alamar blue, SRB staining, and microscopy)
- Data calculated vs 6 DMSO controls as fold-change
- Up and down regulation distinguished
- LELs determined

BioSeek



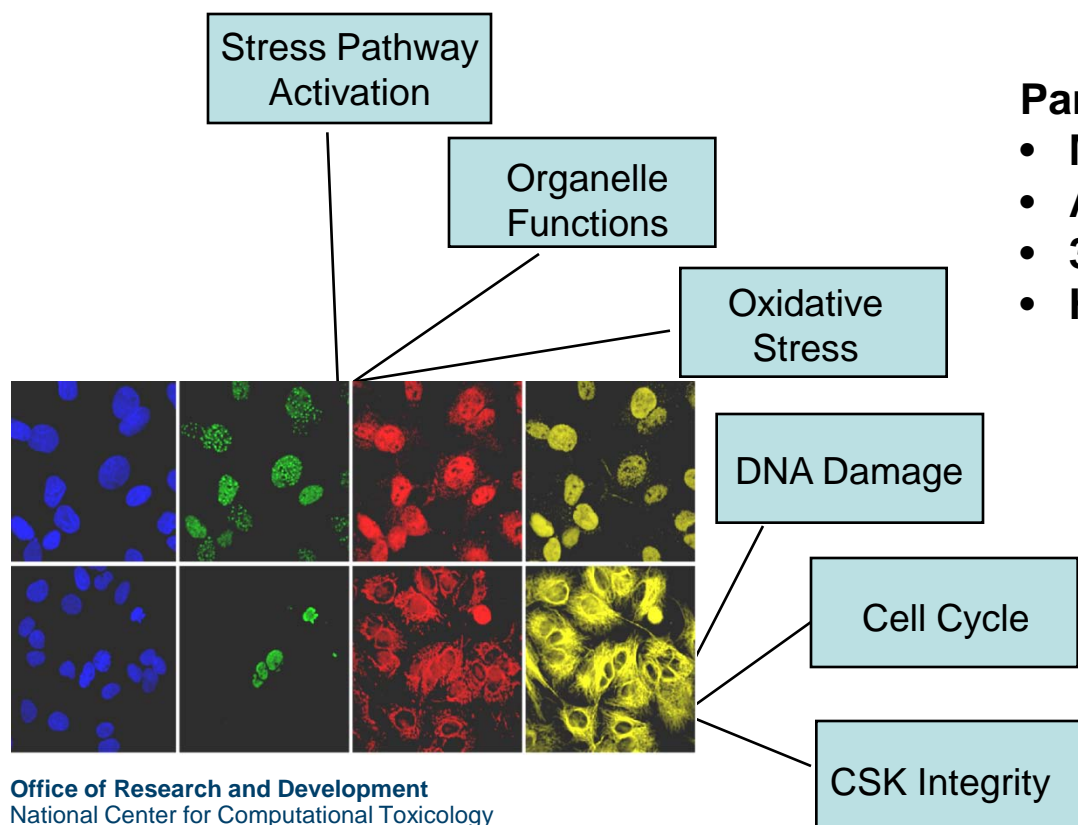
# BioSeek Assays: What is Being Measured?

- Effects of chemicals on signaling pathways in primary human cells
- False positives
  - Cytotoxicity (down-regulation in particular)
  - Statistical vs biological significance
  - Operational
- False negatives
  - Solubility
  - Lack of biotransformation
  - Cytotoxicity
  - Operational



# Cellumen: High-Content Screening of Cellular Phenotypic Toxicity Parameters (Abstract 38)

- Technology: automated fluorescent microscopy
- Objective: Determine effects of chemicals on toxicity biomarkers in a cell culture of HepG2 and primary rat hepatocytes



## Panel 1 design\*:

- Multiple mechanisms of toxicity
- Acute, early & chronic exposure
- 384-well capacity
- HepG2

# Cellumen: CellCiphr™ Cytotoxicity Panel

- 10-point conc-response (200  $\mu$ M-39 nM) in duplicate
- Three time points (1 hr, 24 hr, 72 hr)
- 11 endpoints per assay

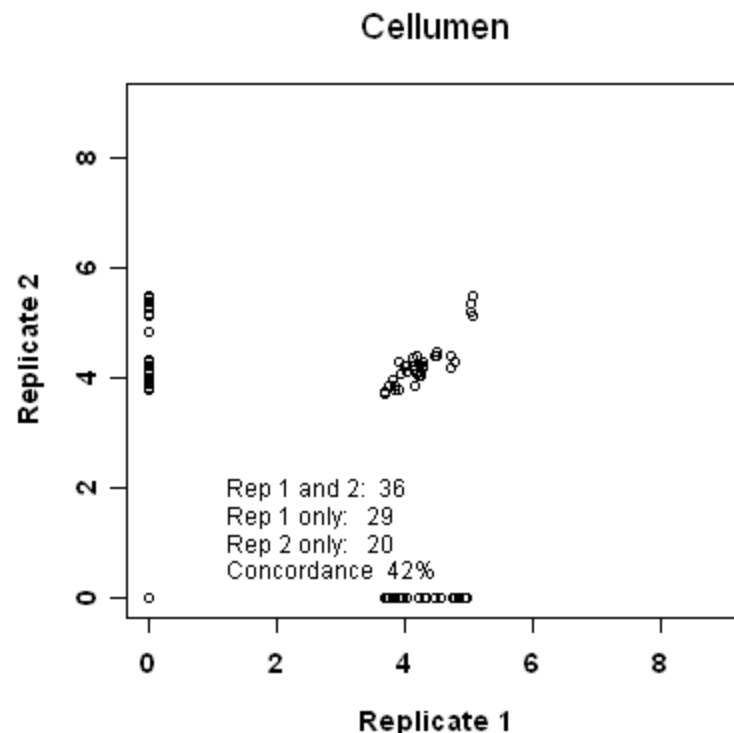
Biomarker	Measurement	Positive Control	Z'
Stress Pathway	Phospho-c-jun	Anisomycin	0.63
Oxidative Stress	Phospho-Histone H2A.X	Camptothecin	0.7
Mitochondrial Function	Mitochondrial membrane potential	CCCP	0.55
Mitochondrial Mass	Mitochondrial mass	CCCP	0.35
Cell Loss	Cell number	Camptothecin	0.56
Cell Cycle	DNA content	Paclitaxel	0.54
DNA Degradation	DNA structure	Paclitaxel	0.6
Nuclear Size	Area of nuclear region	Paclitaxel	0.63
DNA Damage	Detection of p53	Camptothecin	0.43
Mitotic Arrest	Phospho-Histone-H3	Paclitaxel	0.63
Cytoskeletal Integrity	Detection of $\alpha$ -tubulin	Paclitaxel	0.3

# Cellumen: What is Being Measured?

- Cellular toxicity phenotypes in a transformed cell line
- False positives
  - Imaging artifacts
  - Fluorescent compounds
  - Statistical vs biological significance
  - Operational
- False negatives
  - Solubility
  - Cytotoxicity
  - Lack of biotransformation
  - Operational

# Cellumen: Data Calculation

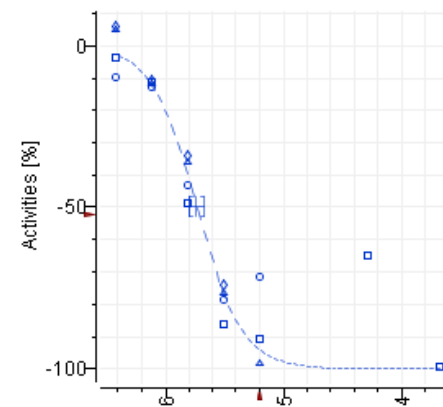
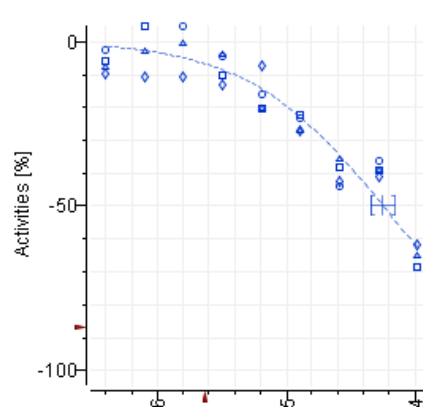
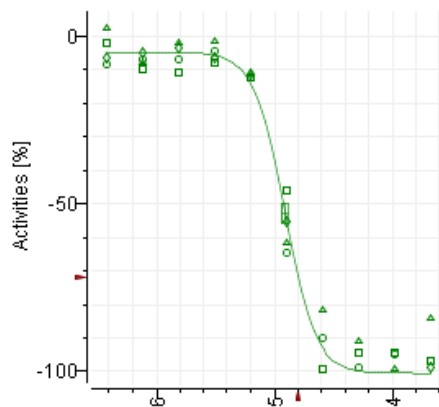
- Data not normalized to controls (with a few exceptions)
- Fit to 3- or 4-parameter Hill equation
- AC50 reported with these rules:
  - For Cell Loss, AC50 is reported as calculated
  - For other endpoints, if AC50 for endpoint is > AC50 for Cell Loss at the corresponding exposure time, AC50 for endpoint is set to 100  $\mu$ M (to account for imaging artifacts of cytotoxicity)
- Issues:
  - Lack of positive controls for all endpoint/time combinations
  - Large differences in maximal response
  - noisy curves due in part to effects of cytotoxicity



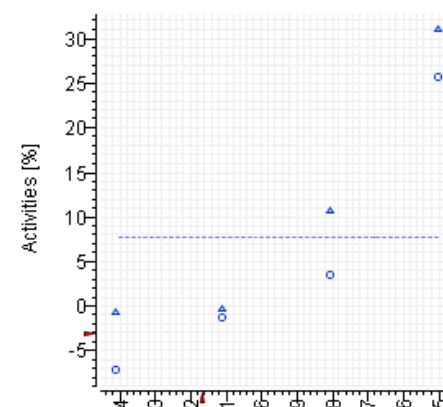
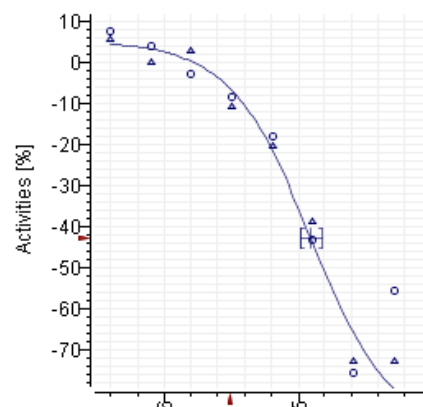
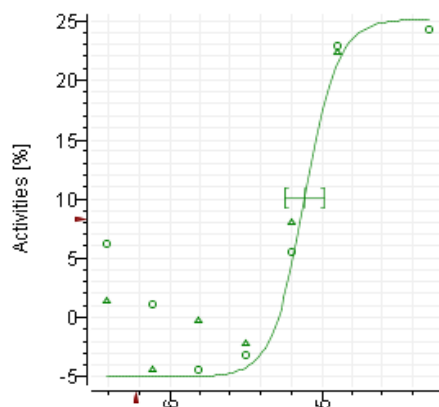


# Cellumen: Data Examples

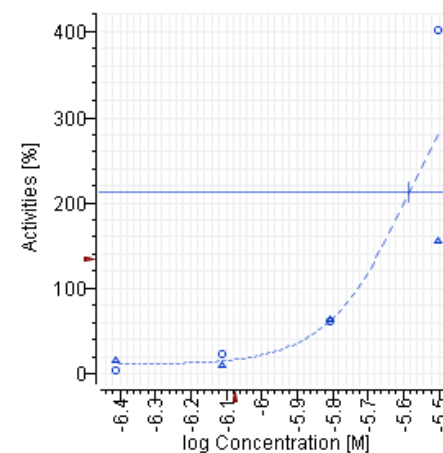
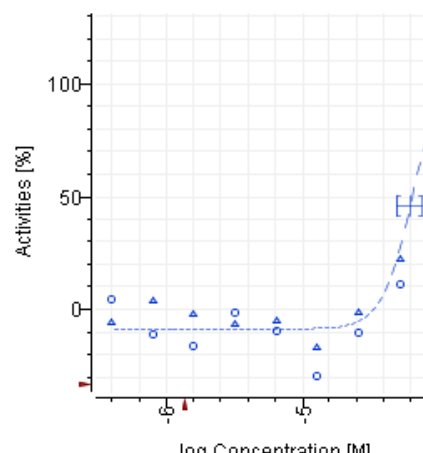
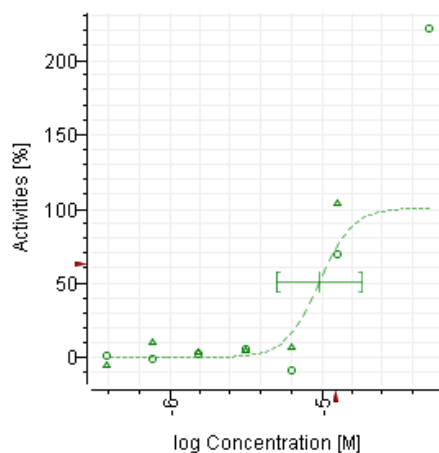
## Cell Loss



## Mitochondrial Membrane Potential

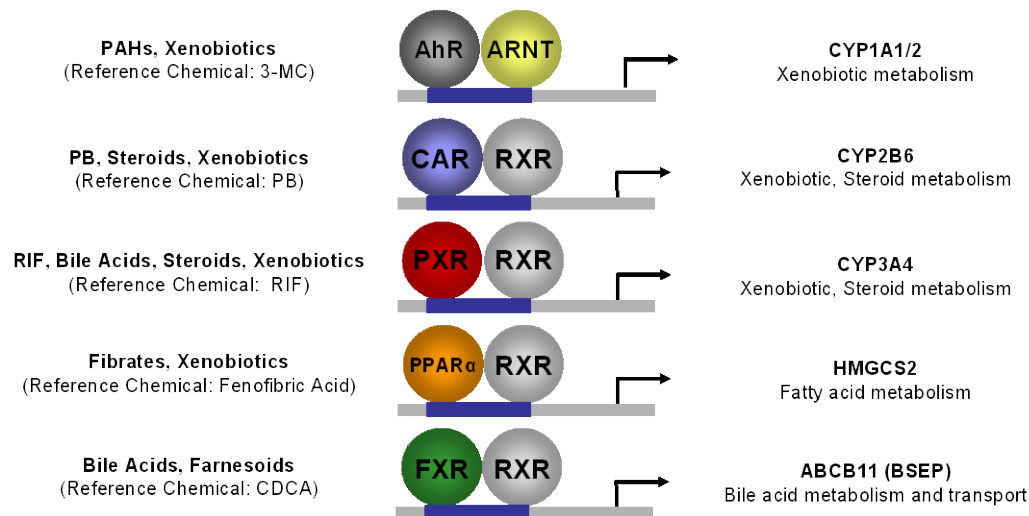


## DNA Damage



# CellzDirect: XME Gene Expression in Primary Human Hepatocytes (Abstract 22)

- Primary human hepatocytes from two donors used
- Cells exposed for 6, 24, and 48 hr; medium/chemical refreshed daily
- Concentrations tested: 40, 4, 0.4, 0.04, and 0.004  $\mu\text{M}$
- 16 Genes measured in multiplexed RNase protection assay (qNPA)
- Genes targeted XME and transporters



## Target Gene Categories

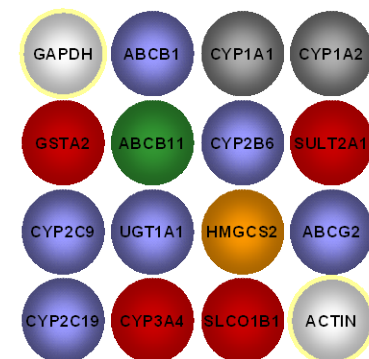
CYP450 (6)

Transporter (4)

Phase II Metabolism (3)

Cholesterol Synthesis (1)

Endogenous control (2)

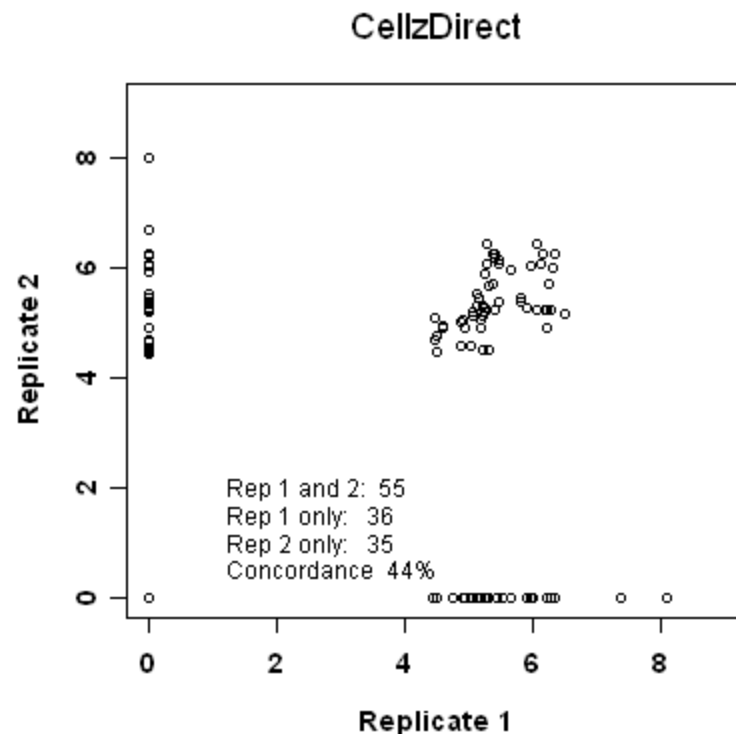


# CellzDirect: What is Being Measured?

- Up/down regulation of mRNA for XME and transporters in primary human hepatocytes
- False positives:
  - General effect of cytotoxicity on transcriptional activity
  - Statistical vs biological significance
  - Operational
- False negatives:
  - Solubility
  - Cytotoxicity
  - Operational
  - Lack of biotransformation
  - Inter-individual donor variation

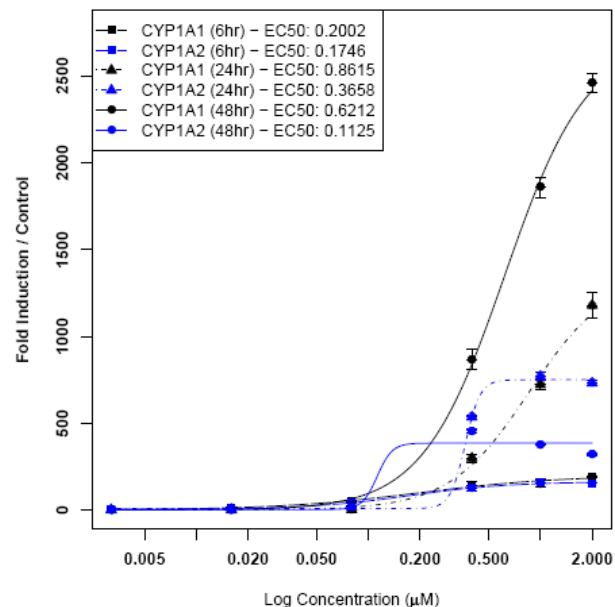
# CellzDirect: Data Calculation

- Normalized to solvent control; expressed as fold-change
- Curves fit to Hill equation
  - Upper and lower limits defined by minimum and maximum responses observed over dataset of a particular gene/donor/time
- LELs determined
- Large variations in maximal responses
  - Biological vs statistical significance
- Measuring endogenous promoter activity reflects complex, multifactorial regulation of gene expression
- 6 hr exposure data not provided due to high variability associated with limited time for gene induction

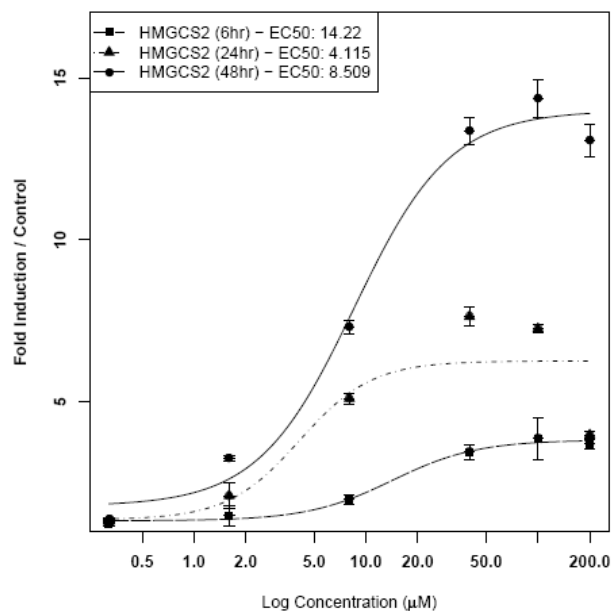


# CellzDirect: Data Examples

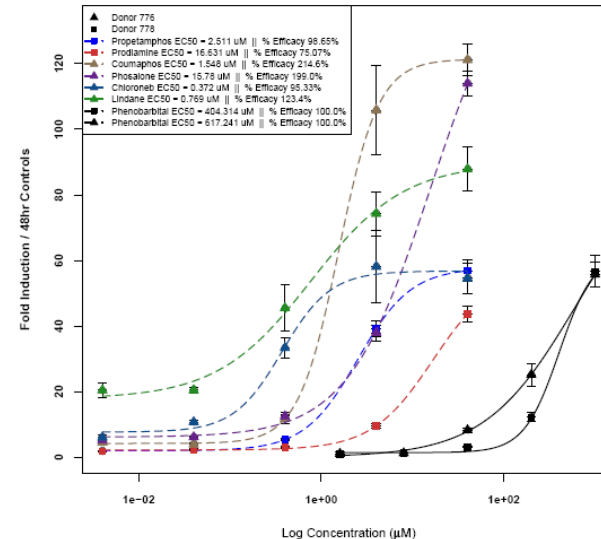
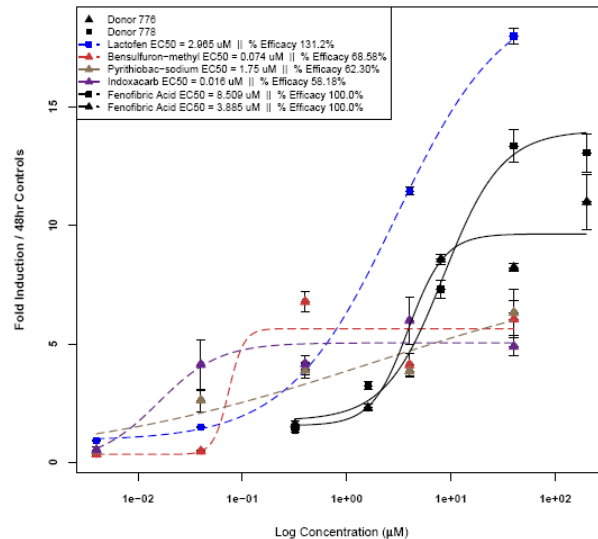
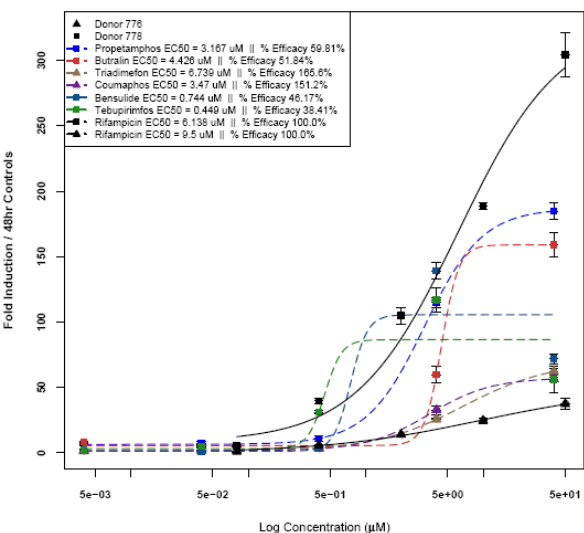
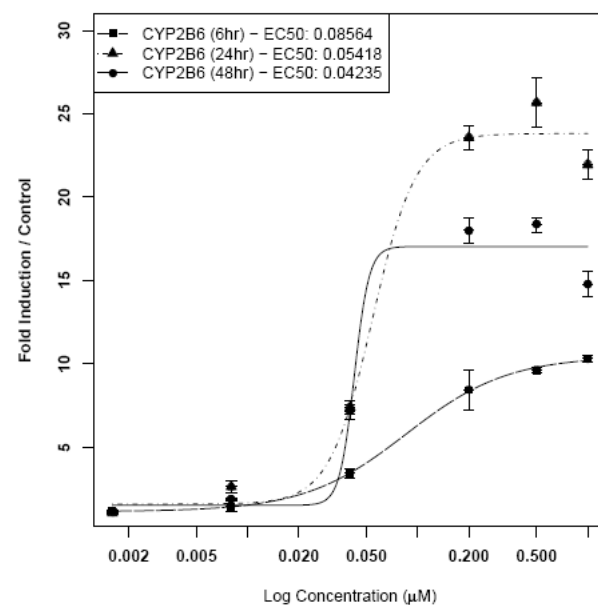
## CYP1A1-AhR



## HMGCS2-PPAR $\alpha$

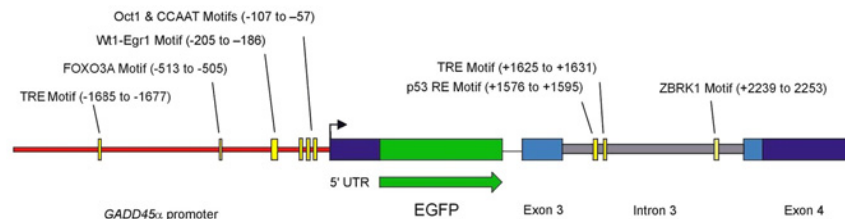
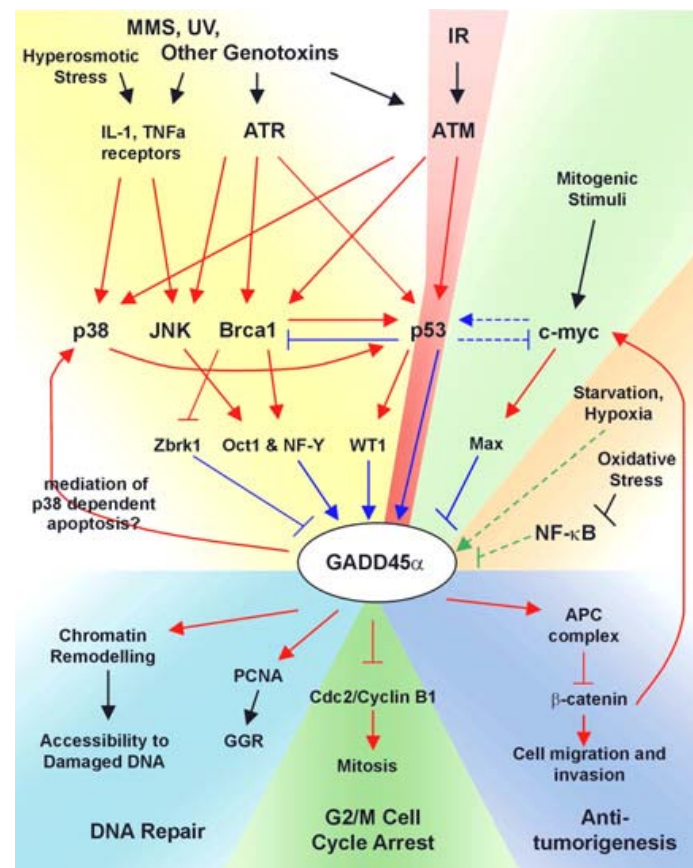


## CYP2B6-CAR



# Gentronix: GADD45a Reporter Gene Assay for DNA Damage (Abstract 41)

- TK6 cell line expressing GFP under control of GADD45a promoter
- Cells exposed at 200, 100, 50  $\mu$ M for 24 and 48 hr
- Cytotox assay to discount artifacts
- Retested at lower conc if cytotoxic

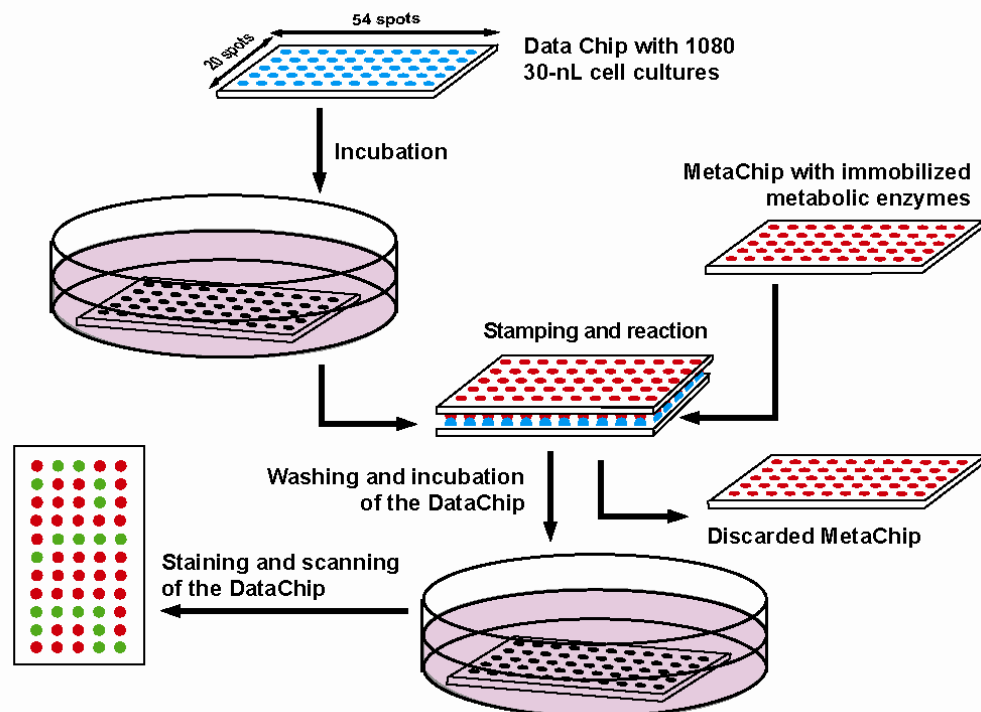


# Gentronix: Data Calculations

- If induction of GFP fluorescence >50%1 test called genotoxic
- If over 2 or 3 concentrations, strongly genotoxic
- LELs calculated
- Replicate analysis: no actives among replicates
- GADD45a promoter activity in transformed cell line
- False positives:
  - Cytotoxicity resulting in general transcriptional activity
  - Cellular stress other than DNA damage
  - Statistical vs biological validation
  - Operational
- False negatives:
  - Solubility
  - Cytotoxicity
  - Lack of biotransformation
  - Operational

# Solidus: Biotransformation Chip and Effect of Cytotoxicity (Abstract 30)

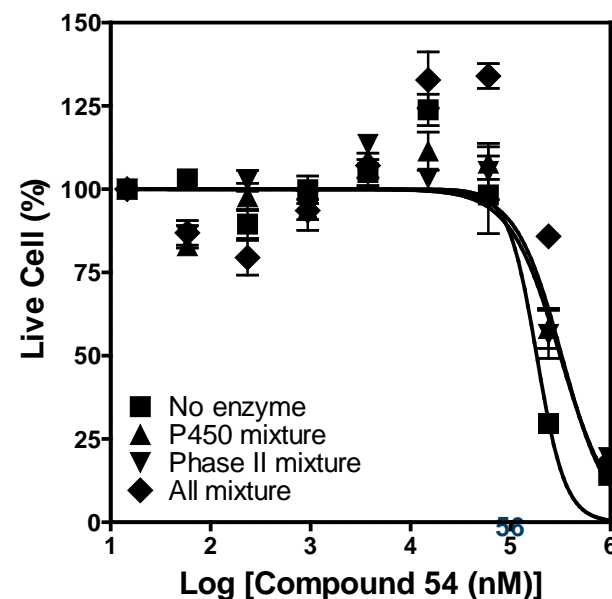
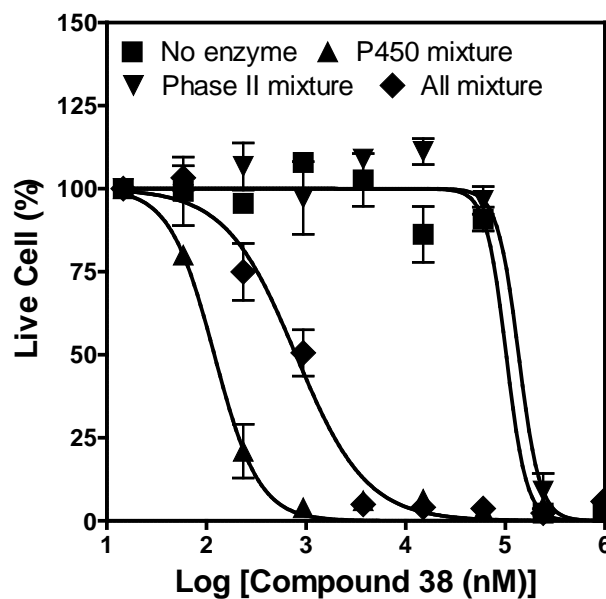
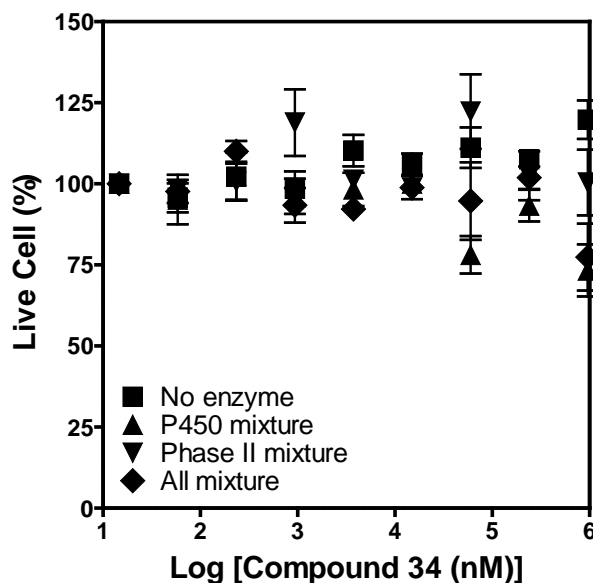
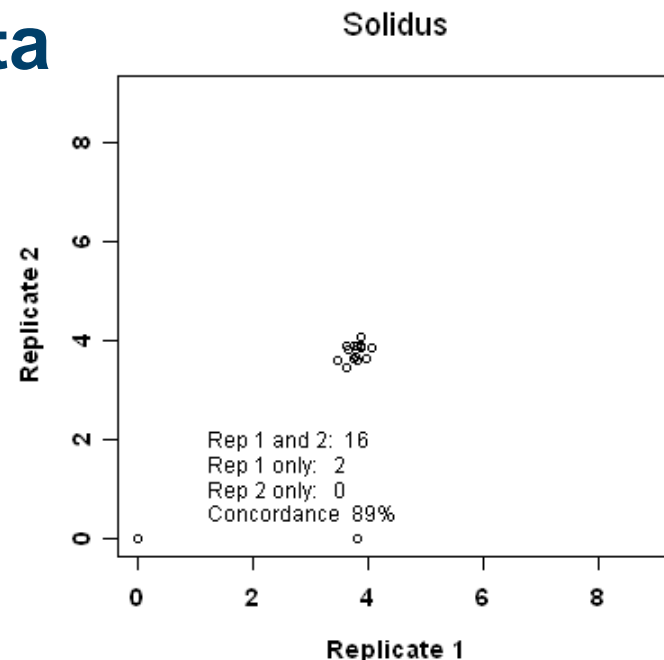
- Alginate-immobilized Phase I and Phase II enzymes
- ToxCast\_320 exposed 6 hr to:
  - Control
  - Ph I
  - Ph II
  - Ph I and Ph II
- 960 uM high conc/4-fold serial dilutions/9 concentrations/5 replicates
- Cytotoxicity in Hep3B measured 48 hr later





# Solidus: Example Data

- Data normalized to control values
- Concentration-response data fit to Hill equation
- LC50 determined for each assay condition



# Solidus: What is Being Measured?

- Effect of Phase I and Phase II enzymes on cytotoxicity activity of chemicals against a transformed cell line
- False positives:
  - Possibly not optimized Phase I and/or Phase II mix
  - Operational
- False negatives
  - Solubility
  - Possibly not optimized Phase I and/or Phase II mix
  - Availability of compound from alginate-immobilized enzyme matrix
  - Operational

# NCGC Reporter Gene Assays

- Nuclear Receptors
  - GAL4 System (ligand detection assay)
  - 11 human receptors
  - 1 rat (PXR)
  - $\beta$ -lactamase reporter gene assays except:
  - PXR assays are luciferase reporter gene assays
- p53 Reporter Gene assay
  - $\beta$ -lactamase reporter gene assay
- Parental cell lines mostly HEK293 (also HeLa and DPX-2)
- 12-15 point concentration-response curves (single replicate)

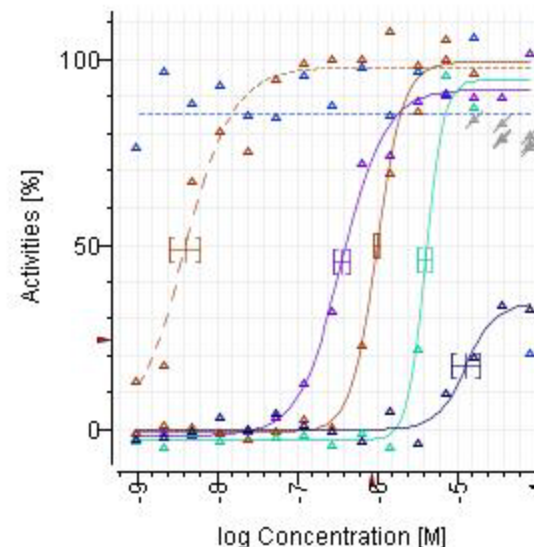


United States  
Environmental Protection  
Agency

# NCGC: Data Calculations

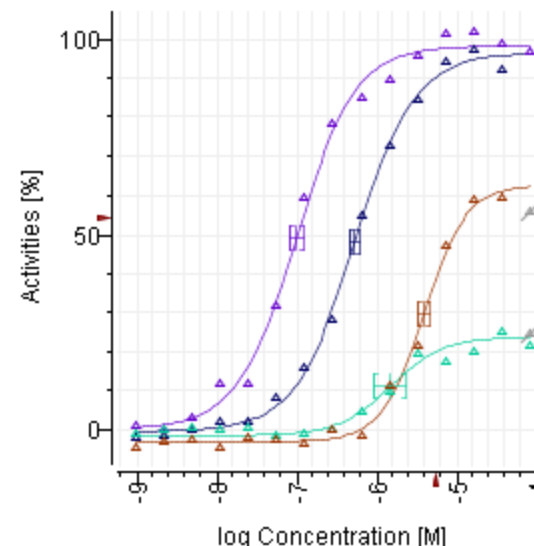
- Data normalized to reference compound effect
- Curves fit to 3- or 4-parameter Hill equation
- Artifacts removed where obvious fluorescence or cytotoxicity detected
- Required at least 25% efficacy of control compound to calculate AC50
- AC50 values provided
- Antagonist format assays challenging due to effects of cytotoxicity
- LXR assay problematic—contaminated with GR reporter line?

ER $\alpha$



■ NCGC00090749-04  
■ NCGC00161666-02

PPAR $\gamma$



■ NCGC00164420-01  
■ NCGC00093991-03  
■ NCGC00164230-01  
■ NCGC00022570-07

# NCGC Assays: What is Being Measured?

- NR assays are ligand-detection assays
- False positives
  - Fluorescent compounds
  - Statistical vs biological significance
  - Gal4\_NR-LBD not physiological
  - Cytotoxicity (antagonist format)
  - Operational
- False negatives
  - Fluorescent compounds
  - Cytotoxicity
  - Gal4\_NR-LBD not physiological
  - Operational
  - Lack of biotransformation

# Additional Data Sets To Be Added Soon:

- NHEERL
  - Zebrafish developmental toxicity (Padilla) Poster
  - Neurite outgrowth and neuronal proliferation (Mundy and Shafer)
  - ES cell differentiation (Hunter)
- Plasma protein binding and hepatocyte clearance (Thomas)
- PPAR $\alpha$  and AhR (NCGC)
- Primary rat hepatocyte HCS Cell Health (Cellumen)